

GenCore version 4.5
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OM nucleic - nucleic search, using sw model

Run on: April 10, 2002, 21:32:13 ; Search time 107.54 Seconds
(without alignments)
2672.499 Million cell updates/sec

Title: US-09-380-276A-1
Perfect score: 1269
Sequence: 1 atgctttaaagtctact.....ggcagcagctgggttccctg 1369

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 351203 seqs, 113238999 residues

Total number of hits satisfying chosen parameters: 702406

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_NA.*
1: /cgn2.6/ptodata/2/ina/5A_COMB.seq.*
2: /cgn2.6/ptodata/2/ina/5B_COMB.seq.*
3: /cgn2.6/ptodata/2/ina/6A_COMB.seq.*
4: /cgn2.6/ptodata/2/ina/6B_COMB.seq.*
5: /cgn2.6/ptodata/2/ina/PCTUS_COMB.seq.*
6: /cgn2.6/ptodata/2/ina/backfiles1.seq.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	435.4	34.3	893	US-09-286-529-8	Sequence 8, Appli
2	314	24.7	623	US-09-286-529-9	Sequence 9, Appli
3	36.4	2.9	1601	US-08-722-001-7	Sequence 7, Appli
4	36.4	2.9	1987	US-08-722-001-26	Sequence 26, Appli
5	36.4	2.9	1997	US-08-722-001-27	Sequence 27, Appli
6	36.4	2.9	2004	US-08-722-001-11	Sequence 11, Appli
7	36.2	2.9	2485	US-08-424-424B-1	Sequence 1, Appli
8	36.2	2.9	2486	PCT-US94-05363A-1	Sequence 1, Appli
9	36	2.8	4136	US-09-103-875-2	Sequence 2, Appli
10	35.6	2.8	1150	US-09-372-934-3	Sequence 3, Appli
11	34.8	2.7	1639	US-08-334-698-5	Sequence 5, Appli
12	34.8	2.7	1639	US-08-228-932-5	Sequence 5, Appli
13	34.8	2.7	1639	US-08-468-939-5	Sequence 5, Appli
14	34.8	2.7	1639	US-08-406-855A-5	Sequence 5, Appli
15	34.8	2.7	1639	US-08-722-190-5	Sequence 5, Appli
16	34.8	2.7	1639	US-08-244-354-5	Sequence 5, Appli
17	34.8	2.7	1639	US-09-206-899-5	Sequence 5, Appli
18	34.8	2.7	1639	PCT-US95-04203-5	Sequence 5, Appli
19	34.6	2.7	5962	US-08-232-463-14	Sequence 14, Appli
20	34.6	2.7	7218	US-08-416-603-11	Sequence 11, Appli
21	33.6	2.6	800	US-08-325-547-9	Sequence 9, Appli
22	33.4	2.6	4360	US-08-200-512-1	Sequence 1, Appli
23	33	2.6	9472	US-08-524-828-2	Sequence 2, Appli
24	32.8	2.6	2230	US-08-975-114A-2	Sequence 1, Appli
25	32.6	2.6	1593	US-08-849-281A-2	Sequence 2, Appli
26	32.6	2.6	1593	US-08-849-281A-2	Sequence 2, Appli
27	32.6	2.6	1593	US-08-849-281A-2	Sequence 2, Appli

28	32.6	2.6	2247	2	US-08-524-828-1	Sequence 1, Appli
29	32.6	2.6	2247	2	US-08-975-114A-1	Sequence 1, Appli
c 30	32.6	2.6	3891	1	US-08-480-604A-27	Sequence 27, Appli
c 31	32.6	2.6	3891	2	US-08-405-496A-27	Sequence 27, Appli
c 32	32.6	2.6	3891	4	US-08-915-136-27	Sequence 27, Appli
33	32.4	2.6	1167	1	US-07-960-985-1	Sequence 1, Appli
34	32.4	2.6	1167	2	US-08-496-671-1	Sequence 1, Appli
35	32.4	2.6	1280	4	US-09-096-776B-4	Sequence 4, Appli
36	32.4	2.6	1491	4	US-09-082-092-9	Sequence 9, Appli
37	32.4	2.6	1524	4	US-08-840-767-3	Sequence 3, Appli
c 38	32.4	2.6	1690	2	US-08-461-812-3	Sequence 3, Appli
39	32.4	2.6	3083	2	US-08-480-994-36	Sequence 36, Appli
40	32.4	2.6	3083	2	US-08-616-844-36	Sequence 36, Appli
41	32.4	2.6	3083	2	US-08-599-654-36	Sequence 36, Appli
42	32.4	2.6	3083	2	US-08-485-573-36	Sequence 36, Appli
43	32.4	2.6	3083	3	US-08-944-868A-36	Sequence 36, Appli
44	32.4	2.6	3083	3	US-08-944-423A-36	Sequence 36, Appli
45	32.4	2.6	3083	3	US-08-925-743-36	Sequence 36, Appli

ALIGNMENTS

RESULT 1
US-09-286-529-8
; Sequence 8, Application US/09286529
; Patent No. 6297367
; GENERAL INFORMATION:
; APPLICANT: Catharine Tribouley
; TITLE OF INVENTION: NEW MEMBERS OF TNF AND TNFR FAMILIES
; FILE REFERENCE: 1408.003/200130.439C1
; CURRENT APPLICATION NUMBER: US/09/286,529
; CURRENT FILING DATE: 1999-04-05
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 8
; LENGTH: 893
; TYPE: DNA
; ORGANISM: human
US-09-286-529-8

Query Match	34.3%;	Score	435.4;	DB	4;	Length	893;
Best Local Similarity	82.1%;	Pred. NO.	1.1e-125;				
Matches	513;	Conservative	0;	Mismatches	111;	Indels	1;
Gaps	1;						
QY	1	atgctttaaagtctactagaacaagaaacatttttcaactcttttagtattacta	60				
Db	55	atggcactcaagtcctactctctacacaggacggtgctctcgtccactctctccta	114				
QY	61	ggctattgtcatgtaaagtgttccctcgaacaggagactgtagacagcaagaattcag	120				
Db	115	ctccacctggcatgtaaagtgttgcgaacccgagattgcaggcagcaggaattcaag	174				
QY	121	gatcgtctggaacatgttccctcgaacacagtggtggccaggcagtgaggtgtctaa	180				
Db	175	gatcgtctggaacatgttccctcgaacacagtggtggccaggcagtgaggtgtctaa	234				
QY	181	gaatgtggtctcgctcatgtggagagatgcacagtgtgaacgtcccggtcgcacaggttc	240				
Db	235	gaatgtggtctcgctcatgtggagagatgcacagtgtgaacgtcccggtcgcacaggttc	294				
QY	241	aaggagactggggtcttcagaaatgcagccctgtctgactgcgcagtggtgaaaccgc	300				
Db	295	aaggagactggggtcttcagaaatgcagccctgtgactgcgcagtggtgaaaccgc	354				
QY	301	tttcagaagcaaatgtttcagccacacagtgatgcacatctcggggagactgcttgcagg	360				
Db	355	tttcagaagcaaatgtttcagccacacagtgatgcacatctcggggagactgcttgcagg	414				
QY	361	ttttatagaagacgaaactgtcgtcttcaagacatgagtgatgtccttgccttgcagac	420				
Db	415	ttttacccggaagaccacacagtggtgttttcaagacatgagtgatgtccttgccttgcagac	474				

GENERAL INFORMATION:
APPLICANT: Thompson, Wayne J.
TITLE OF INVENTION: ALPHAIC ADRENERGIC RECEPTOR ANTAGONISTS
NUMBER OF SEQUENCES: 35
CORRESPONDENCE ADDRESS:
APPLICANT: Nerenberg, Jennie B.
APPLICANT: Lee, Hee-Yoon
APPLICANT: Bell, Ian M.
STREET: 126 Lincoln Avenue
CITY: Rahway
STATE: New Jersey
COUNTRY: United States of America
ZIP: 07065
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/722,001
FILING DATE:
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/229,276
FILING DATE: 14-APR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Appollina, Mary A.
REGISTRATION NUMBER: 34,087
REFERENCE/DOCKET NUMBER: 19169Y
TELECOMMUNICATION INFORMATION:
TELEPHONE: (908)594-3462
TELEFAX: (908)594-4720
TELEX: 138825
INFORMATION FOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
LENGTH: 1987 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
TOPOLOGY: both
MOLECULE TYPE: cDNA
HYPOTHETICAL: NO
ANTI-SENSE: NO
US-08-722-001-26

Query Match 2.9%; Score 36.4; DB 1; Length 1987;
Best Local Similarity 52.7%; Pred. No. 0.27;
Matches 79; Conservative 0; Mismatches 71; Indels 0; Gaps 0;
QY 474 cgcgtccacggtccagccacgggacacggcgtggtcggtatctgcagcgctct 533
DB 1112 CGAGGACGAGACCATCTGCCAGATCAACGAGGAGCGGGCTACGTGCTCTTCGCGCTCT 1171
QY 534 ggccaccgtctgtggtcggtcgtatctgtggtcatttgaagagacagttat 593
DB 1172 GGGCTCTTCTACCTGCTCTGGCCATCATCTGCTGATGTTACTGCGGCTCTACGTGCT 1231
QY 594 ggagagaaacccagctggtctcgtcggtc 623
DB 1232 GGCCAGAGAGGAGAGCGGGGCTCAAGTC 1261

RESULT 5
US-08-722-001-27
Sequence 27, Application US/08722001
Patent No. 5760054
GENERAL INFORMATION:
APPLICANT: Thompson, Wayne J.
APPLICANT: Huff, Joel R.
APPLICANT: Nerenberg, Jennie B.
APPLICANT: Lee, Hee-Yoon

APPLICANT: Bell, Ian M.
TITLE OF INVENTION: ALPHAIC ADRENERGIC RECEPTOR ANTAGONISTS
NUMBER OF SEQUENCES: 35
CORRESPONDENCE ADDRESS:
APPLICANT: Merck & Co., Inc.
STREET: 126 Lincoln Avenue
CITY: Rahway
STATE: New Jersey
COUNTRY: United States of America
ZIP: 07065
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/722,001
FILING DATE:
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/229,276
FILING DATE: 14-APR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Appollina, Mary A.
REGISTRATION NUMBER: 34,087
REFERENCE/DOCKET NUMBER: 19169Y
TELECOMMUNICATION INFORMATION:
TELEPHONE: (908)594-3462
TELEFAX: (908)594-4720
TELEX: 138825
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
LENGTH: 1997 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
TOPOLOGY: both
MOLECULE TYPE: cDNA
HYPOTHETICAL: NO
ANTI-SENSE: NO
US-08-722-001-27

Query Match 2.9%; Score 36.4; DB 1; Length 1997;
Best Local Similarity 52.7%; Pred. No. 0.27;
Matches 79; Conservative 0; Mismatches 71; Indels 0; Gaps 0;
QY 474 cgcgtccacggtccagccacgggacacggcgtggtcggtatctgcagcgctct 533
DB 1106 CGAGGACGAGACCATCTGCCAGATCAACGAGGAGCGGGCTACGTGCTCTTCGCGCTCT 1165
QY 534 ggccaccgtctgtggtcggtcgtatctgtggtcatttgaagagacagttat 593
DB 1166 GGGCTCTTCTACCTGCTCTGGCCATCATCTGCTGATGTTACTGCGGCTCTACGTGCT 1225
QY 594 ggagagaaacccagctggtctcgtcggtc 623
DB 1226 GGCCAGAGAGGAGAGCGGGGCTCAAGTC 1255

RESULT 6
US-08-722-001-11
Sequence 11, Application US/08722001
Patent No. 5760054
GENERAL INFORMATION:
APPLICANT: Thompson, Wayne J.
APPLICANT: Huff, Joel R.
APPLICANT: Nerenberg, Jennie B.
APPLICANT: Lee, Hee-Yoon
APPLICANT: Bell, Ian M.
TITLE OF INVENTION: ALPHAIC ADRENERGIC RECEPTOR ANTAGONISTS
NUMBER OF SEQUENCES: 35
CORRESPONDENCE ADDRESS:
APPLICANT: Merck & Co., Inc.

REFERENCE/DOCKET NUMBER: 325800-118
TELEPHONE: 201-994-1700
TELEFAX: 201-994-1744
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 2486 BASE PAIRS
TYPE: NUCLEIC ACID
STRANDEDNESS: SINGLE
TOPOLOGY: LINEAR
MOLECULE TYPE: CDNA
PCT-US94-05363A-1

Query Match 2.9%; Score 36.2; DB 5; Length 2486;

Best Local Similarity 56.2%; Pred. No. 0.36;
Matches 68; Conservative 0; Mismatches 53; Indels 0; Gaps 0;

QY 453 caaggtcaacctgtgaagatcgctccacgctccacgacccacgggacacgctgctgc 512

DB 2024 CAGCATCATCCAGCTGGGGTACGCCGCCCGCTACAGCGCTGGATCAAGGAGGAGC 2083

QY 513 tcccttattctcagcgctctgacacgctcctgctgctgctgctcctcctctgtgtcat 572

DB 2084 TCGCGAGCGCTACCTGTATTTCCTCAACTGGCCATGGCCTCTCTGATCACCTCATCGT 2143

QY 573 c 573

DB 2144 C 2144

RESULT 9

US-09-103-875-2/c
Sequence 2, Application US/09103875A
Patent No. 6221849

GENERAL INFORMATION:
APPLICANT: Szyf, Moshe

APPLICANT: Bigey, Pascal

APPLICANT: Ramchandani, Shyam

TITLE OF INVENTION: DNA METHYLTRANSFERASE GENOMIC SEQUENCES AND ANTISENSE

TITLE OF INVENTION: OLIGONUCLEOTIDES

FILE REFERENCE: 106101.194

CURRENT APPLICATION NUMBER: US/09/103,875A

CURRENT FILING DATE: 1998-06-24

EARLIER APPLICATION NUMBER: 60/069,865

EARLIER FILING DATE: 1997-12-17

EARLIER APPLICATION NUMBER: 08/866,340

EARLIER FILING DATE: 1997-05-30

NUMBER OF SEQ ID NOS: 138

SOFTWARE: PatentIn Ver. 2.0

SEQ ID NO 2

LENGTH: 4136

TYPE: DNA

ORGANISM: Homo sapiens

US-09-103-875-2

Query Match 2.8%; Score 36; DB 4; Length 4136;

Best Local Similarity 49.5%; Pred. No. 0.56;
Matches 93; Conservative 0; Mismatches 95; Indels 0; Gaps 0;

QY 689 cccacagacgctgctgcccagtcgcccgtgactcagtcagacacctgcccggcggtgcgt 748

DB 3917 CCCACCCAGCGCCCTGCTGCTCCCTGTGAGTCGTGTCCCTCCCATGTGTACTACCGCC 3858

QY 749 tgcctccatctgctgtgagagcctgacgcccacccacccgacgactcttgggtg 808

DB 3857 TCGCGACATCGTCGGCAGCAGATGCGGGACGGCCAGTGTGGCAGCCCGGGCTGGGG 3798

QY 809 ggggtcattctgagccagctcttcaggcaagaacacgagccacgcccagcgggagatggtgc 868

DB 3797 CGGTAGCGCGGCATCTCGGAGGCTTCAGCAGACCGCGGGCGGCGAGCGCGGCCCC 3738

QY 869 cgaatttc 876
DB 3737 GCGTTTTC 3730

RESULT 10

US-09-372-934-3

Sequence 3, Application US/09372934

Patent No. 6248579

GENERAL INFORMATION:

APPLICANT: Stutzman-Engwall, Kim J.

APPLICANT: McArthur, Ramish

APPLICANT: Kato, Yoshihiro

TITLE OF INVENTION: STREPTOMYCES AVERMITILIS GENE DIRECTING THE RATIO OF

FILE REFERENCE: PC10649

CURRENT APPLICATION NUMBER: US/09/372,934

CURRENT FILING DATE: 1999-08-12

EARLIER APPLICATION NUMBER: 60/074,636

EARLIER FILING DATE: 1998-02-13

EARLIER APPLICATION NUMBER: PCT/IB99/00130

EARLIER FILING DATE: 1999-01-25

NUMBER OF SEQ ID NOS: 25

SOFTWARE: PatentIn Ver. 2.0

SEQ ID NO 3

LENGTH: 1150

TYPE: DNA

ORGANISM: Streptomyces hygroscopicus

FEATURE:

NAME/KEY: CDS

LOCATION: (58)..(990)

US-09-372-934-3

Query Match

Best Local Similarity 2.8%; Score 35.6; DB 4; Length 1150;

Matches 65; Conservative 0; Mismatches 49; Indels 0; Gaps 0;

QY 448 gccagcaagtcacacctgctgaagatcgctccacgctccacgctccacgagggacacgcg 507

DB 142 gccagacgctctaccgcatcagagagcgctcccgccagggcggtgggactcggag 201

QY 508 ctggctgctgcttctgagcgctctgcccacgctgctgctgctgctgctgctc 561

DB 202 cggatcgccgctgctgctgctgctgctgctgctgctgctgctgctgctgctc 255

RESULT 11

US-08-334-698-5

Sequence 5, Application US/08334698

Patent No. 5556753

GENERAL INFORMATION:

APPLICANT: Jonathan A. Bard et al.

TITLE OF INVENTION: DNA Encoding Human Alpha 1 Adrenergic

TITLE OF INVENTION: Receptors and Uses Thereof

NUMBER OF SEQUENCES: 6

CORRESPONDENCE ADDRESS:

ADDRESSEE: COOPER & DUNHAM

STREET: 30 Rockefeller Plaza

CITY: New York

STATE: New York

COUNTRY: U.S.A.

ZIP: 10112

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.24

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/334,698

FILING DATE:

CLASSIFICATION: 435

PRIOR APPLICATION DATA:

RESULT 12
US-08-228-932-5
: Sequence 5, Application US/08228932
: Patent No. 5578611
: GENERAL INFORMATION:
: APPLICANT: Charles Gluchowski, Carlos C. Forray, George Chiu,
: APPLICANT: Theresa A. Brancheck, John M. Wetzel and Paul R. Hartig
: TITLE OF INVENTION: USE OF ALPHA-1C SPECIFIC COMPOUNDS TO TREAT BENIGN
: TITLE OF INVENTION: PROSTATIC HYPERPLASIA
: NUMBER OF SEQUENCES: 6
: CORRESPONDENCE ADDRESS:
: ADDRESSEE: COOPER & DUNHAM
: STREET: 30 Rockefeller Plaza
: CITY: New York
: STATE: New York
: COUNTRY: U.S.A.
: ZIP: 10112
: COMPUTER READABLE FORM:
: MEDIUM TYPE: Floppy disk
: COMPUTER: IBM PC compatible
: OPERATING SYSTEM: PC-DOS/MS-DOS
: SOFTWARE: PatentIn Release #1.24
: CURRENT APPLICATION DATA:
: APPLICATION NUMBER: US/08/228,932
: FILING DATE: 13-APR-1994
: CLASSIFICATION: 514
: ATTORNEY/AGENT INFORMATION:
: NAME: White, John P.
: REGISTRATION NUMBER: 28,678
: REFERENCE/DOCKET NUMBER: 41878-B/JPW/TEP

RESULT 13
US-08-468-939-5
; Sequence 5, Application US/08468939
; Patent No. 5714381
; GENERAL INFORMATION:
; APPLICANT: Jonathan A. Bard et al.
; TITLE OF INVENTION: DNA Encoding Human Alpha 1 Adrennergic
; TITLE OF INVENTION: Receptors and Uses Thereof
; NUMBER OF SEQUENCES: 6
; CORRESPONDENCE ADDRESS:
; ADDRESS: COOPER & DUNHAM LLP
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10036
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.24
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/468.939
; FILING DATE:
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: White, John P.
; REGISTRATION NUMBER: 28,678
; REFERENCE/DOCKET NUMBER: 41337-1B
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212) 278-0400
; TELEFAX: (212) 391-0526
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1639 base pairs
; TYPE: nucleic acid

	Matches	78;	Conservative	0;	Mismatches	72;	Indels	0;	Gaps	0;
QY 474	cgcgtccacg	ggcctccag	ccacggg	gacacg	gcgtg	gctg	cggtt	atct	gcag	gctct 533
Db 635	CGAGGACG	AGACCAT	CTGCC	ATCAAC	GAGAG	CCGGG	CTACG	TCTCT	CACG	CGCT 594
QY 534	ggccaccg	tcctg	ctg	ggcc	ctg	ctc	ctc	ctg	tg	tc
Db 695	GGGCTCCT	TCTAC	CTGC	CTTG	CCAT	CATC	CTTG	GTCA	TGTAC	TGCGGCTCTACG
QY 594	ggagaaga	aaaccc	agctg	gtg	ctc	tc	tc	tc	gc	ggtc 623
Db 755	GGCCAA	GAGAG	AGAG	CCGG	GGCT	CAAG	TC			784

Search completed: April 11, 2002, 00:22:56
 Job time: 10243 sec

GenCore version 4.5
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OM nucleic - nucleic search, using sw model

Run on: April 11, 2002, 00:22:56 ; Search time 107.54 Seconds
(without alignments)
3588.604 Million cell updates/sec

Title: US-09-380-276A-2
Perfect score: 1704
Sequence: 1 ggggaacgtagaactctccaa.....gaccagagtatacttttc 1704

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 351203 seqs, 113238999 residues

Total number of hits satisfying chosen parameters: 702406

Minimum DB seq length: 0
Maximum DB seq length: 2000000000
Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_NA.*
1: /cgn2_6/ptodata/2/ina/5A_COMB.seq.*
2: /cgn2_6/ptodata/2/ina/5B_COMB.seq.*
3: /cgn2_6/ptodata/2/ina/6A_COMB.seq.*
4: /cgn2_6/ptodata/2/ina/6B_COMB.seq.*
5: /cgn2_6/ptodata/2/ina/PCTUS_COMB.seq.*
6: /cgn2_6/ptodata/2/ina/backfiles1.seq.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	447.2	26.2	893	4	US-09-386-529-8
2	326.6	19.2	623	4	US-09-286-529-9
3	36.4	2.1	1601	1	US-08-722-001-7
4	36.4	2.1	1987	1	US-08-722-001-26
5	36.4	2.1	1997	1	US-08-722-001-27
6	36.4	2.1	2004	1	US-08-722-001-11
7	36.2	2.1	2485	1	US-08-424-424B-1
8	36.2	2.1	2486	5	PCT-US94-05363A-1
9	36	2.1	4136	4	US-09-103-875-2
10	35.6	2.1	1150	4	US-09-372-934-3
11	35.4	2.1	7218	1	US-08-232-463-14
12	34.8	2.0	1639	1	US-08-334-698-5
13	34.8	2.0	1639	1	US-08-228-932-5
14	34.8	2.0	1639	1	US-08-468-939-5
15	34.8	2.0	1639	2	US-08-406-855A-5
16	34.8	2.0	1639	2	US-08-722-190-5
17	34.8	2.0	1639	3	US-08-244-354-5
18	34.8	2.0	1639	3	US-09-206-899-5
19	34.8	2.0	1639	5	PCT-US95-04203-5
20	34.6	2.0	5962	6	5386025-5
21	33.6	2.0	800	2	US-08-416-603-11
22	33.4	2.0	4360	1	US-08-470-350B-1
23	33	1.9	9472	1	US-08-325-547-9
24	32.8	1.9	2230	1	US-08-200-512-1
25	32.6	1.9	1593	2	US-08-524-828-2
26	32.6	1.9	1593	2	US-08-975-114A-2
27	32.6	1.9	1593	3	US-08-849-281A-2

28	32.6	1.9	2247	2	US-08-524-828-1	Sequence 1, Appli
29	32.6	1.9	2247	1	US-08-975-114A-1	Sequence 1, Appli
C 30	32.6	1.9	3891	1	US-08-480-604A-27	Sequence 27, Appli
C 31	32.6	1.9	3891	2	US-08-405-496A-27	Sequence 27, Appli
C 32	32.6	1.9	3891	4	US-08-915-136-27	Sequence 27, Appli
33	32.4	1.9	1167	1	US-07-960-985-1	Sequence 1, Appli
34	32.4	1.9	1167	2	US-08-496-671-1	Sequence 1, Appli
35	32.4	1.9	1280	4	US-09-096-776B-4	Sequence 4, Appli
36	32.4	1.9	1491	4	US-09-082-092-9	Sequence 9, Appli
37	32.4	1.9	1524	4	US-08-840-767-3	Sequence 3, Appli
C 38	32.4	1.9	1690	2	US-08-461-812-3	Sequence 3, Appli
39	32.4	1.9	3083	2	US-08-480-994-36	Sequence 36, Appli
40	32.4	1.9	3083	2	US-08-616-844-36	Sequence 36, Appli
41	32.4	1.9	3083	2	US-08-599-654-36	Sequence 36, Appli
42	32.4	1.9	3083	2	US-08-485-573-36	Sequence 36, Appli
43	32.4	1.9	3083	3	US-08-944-868A-36	Sequence 36, Appli
44	32.4	1.9	3083	3	US-08-944-423A-36	Sequence 36, Appli
45	32.4	1.9	3083	3	US-08-925-743-36	Sequence 36, Appli

ALIGNMENTS

RESULT 1
US-09-286-529-8
; Sequence 8, Application US/09286529
; Patent No. 6297367
; GENERAL INFORMATION:
; APPLICANT: Catherine Tribouley
; TITLE OF INVENTION: NEW MEMBERS OF TNF AND TNFR FAMILIES
; FILE REFERENCE: 1408.003/200130.439C1
; CURRENT APPLICATION NUMBER: US/09/286,529
; CURRENT FILING DATE: 1999-04-05
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 8
; LENGTH: 893
; TYPE: DNA
; ORGANISM: human
US-09-286-529-8

Query Match	26.2%	Score 447.2;	DB 4;	Length 893;
Best Local Similarity	81.6%	Pred. No. 1.4e-127;		
Matches 529;	Conservative	0;	Mismatches 118;	Indels 1;
Gaps	1;			
Qy	22	aataatacatttgataagaagatggcttttaaaagtctactagaacaagaaaacgt	81	
Db	32	aataaacacgtttgtgagagccatggcactcaagggtctcctctacacagggacgtgc	91	
Qy	82	tttcaactcttttagtattactaggctattgtcatgtaaaagtgaactgtgaacacagag	141	
Db	92	tcttcgtgcattctcttctactccactggcatgtaaaagtgaactgtgcaaacccgag	151	
Qy	142	actgtagacgaagaattcaggatcggtctggaactgtgtccctgcacaccagtgtg	201	
Db	152	attgcaggcagcaggaattcaagatcgatctggaactgtgtcctctgcaacagtgcg	211	
Qy	202	ggccagcatgagattcttaagaattggtctcgctgctggaaggagcacagtgtg	261	
Db	212	gacctgcatggaattgtcccaaggaatggcttcggtcgtatggggaggtcacagtgtg	271	
Qy	262	tgacgtgccggtctcacagggttcgaaggaggtggggttcctccagaaatgcaagccctgc	321	
Db	272	tgccctgcaggccgacccggttcgaaggaggtgggtttccagaagtgttaagccatgtg	331	
Qy	322	tggactgcgagtggtgaaccgctttcagaaggcgaattgttcagccaccagtgtgcga	381	
Db	332	cggactgtgcgtgtggaaccgctttcagaaggcgaattgttcagccaccagtgtgtg	391	
Qy	382	tctgcgggactgttccagagatttatagaagaacgaaactgtggtcttcaagaca	441	
Db	392	tctgcgggactgtcctccagggtttttaccoggaagacaaactgtgtgttttcaagaca	451	

RESULT 5
US-08-722-001-27
; Sequence 27, Application US/08722001
; Patent No. 5760054
; GENERAL INFORMATION:

RESULT 6
US-08-722-001-11
; Sequence 11, Application US/08722001
; Patent No. 5760054
; GENERAL INFORMATION:
; APPLICANT: Thompson, Wayne J.
; APPLICANT: Huff, Joel R.
; APPLICANT: Nerenberg, Jennie B.
; APPLICANT: Lee, Hee-Yoon
; APPLICANT: Bell, Tan M.

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/232,463
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/935,313
FILING DATE:
APPLICATION NUMBER: EP 91 114 300.6
FILING DATE: 26-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 30472/114 IMMU
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)836-9300
TELEFAX: (703)683-4109
TELEX: 899149
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 7218 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
IMMEDIATE SOURCE:
CLONE: pTZgpt-F1s
US-08-232-463-14

Query Match 2.1%; Score 35.4; DB 1; Length 7218;
Best Local Similarity 6.4%; Pred. No. 2.1;
Matches 27; Conservative 203; Mismatches 189; Indels 0; Gaps 0;

QY 11 aactctcaacaataacattatagaagaagatgctttaagaagtctactagaaca 70
DB 1481 AATACCTATCTATCACTAGTAAAGAGATAGAGAAATTTGGTACRRRRRRRRRR 1422
QY 71 agagaaacgttttcaactcttttagtattactaggtctattgtcatgtaaaagtacttg 130
DB 1421 RRR 1362
QY 131 tgaacagagagacttagacacagaaatcaggatcggtctggaactgtgtccctg 190
DB 1361 RRR 1302
QY 191 caacagtggtgagcagcatgagttgtctaagaatgtgcttgcgtatggaggga 250
DB 1301 RRR 1242
QY 251 tgcacagtgtgtgacgtgcccgtgcacaggttcaaggaggagactggggttccagaaatg 310
DB 1241 RRR 1182
QY 311 caagccctgtctgactgcagtggtggaacgcttccagaaggcaaatgttccagccac 370
DB 1181 RRR 1122
QY 371 cagtgtgccatctgaggagactgttccaggaatttatagaagagaaactgtctg 429
DB 1121 RRR 1063

RESULT 12
US-08-334-698-5
Sequence 5, Application US/08334598
Patent No. 5556753
GENERAL INFORMATION:
APPLICANT: Jonathan A. Bard et al.
TITLE OF INVENTION: DNA Encoding Human Alpha 1 Adrenergic
RECEPTORS AND USES THEREOF
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSEE: COOPER & DUNHAM
STREET: 30 Rockefeller Plaza

CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10112
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.24
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/334,698
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/952,798
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 376901
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 977-9550
TELEFAX: (212) 664-0525
TELEX: (212) 42523 COOP UI
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1639 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: unknown
MOLECULE TYPE: DNA (genomic)
HYPOTHETICAL: N
ANTI-SENSE: N
FEATURE:
NAME/KEY: CDS
LOCATION: 126..1523
OTHER INFORMATION:
US-08-334-698-5

Query Match 2.0%; Score 34.8; DB 1; Length 1639;
Best Local Similarity 52.0%; Pred. No. 1.3;
Matches 78; Conservative 0; Mismatches 72; Indels 0; Gaps 0;

QY 518 cgcgtccacggcctccagccacgggacagcggtggtgcgttatctgcagcgctct 577
DB 635 CGAGGACGAGACCATCTGCCAGATCAACGAGGCGGGCTACTGCTCTCTCTCAGCGCT .694
QY 578 ggcacccgtcctgtgcccctctcatcctctgtgtcatctattgtaagagacagttat 637
DB 695 GGGCTCTTCTACCTGCTCTGCGCATCATCTCTGCTCATCTACTGCCGCTCTACGTGCT 754
QY 638 ggagaagaaacccagctggtctctctcggtc 667
DB 755 GGCCAAGAGGAGAGCGCGGGCTCAAGTC 784

RESULT 13
US-08-228-932-5
Sequence 5, Application US/08228932
Patent No. 5578611
GENERAL INFORMATION:
APPLICANT: Charles Gluchowski, Carlos C. Forray, George Chiu,
APPLICANT: Theresa A. Brancheck, John M. Wetzel and Paul R. Hartig
TITLE OF INVENTION: USE OF ALPHA-1C SPECIFIC COMPOUNDS TO TREAT BENIGN
PROSTATIC HYPERPLASIA
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSEE: COOPER & DUNHAM
STREET: 30 Rockefeller Plaza
CITY: New York
STATE: New York
COUNTRY: U.S.A.

```

Query Match      2.0%; Score 34.8; DB 1; Length 1639;
Best Local Similarity 52.0%; Pred. No. 1.3;
Matches 78; Conservative 0; Mismatches 72; Indels 0; Gaps 0;

QY 518 cgcgtccacggcctccagccacgggacacggcgctggctgcggttatctgcagcgtct 577

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QY	578	ggccacccgctctgctgagccctgctcaatcctctgtgtctcatctatctgttaagagaacaggtttat	637
Db	695	ggcgtctcttcttacctgcctctctggccatcatctgtctcatctgtctacgtggt	754
QY	638	ggagaagaaccccgctggtctctcggtc	667
Db	755	ggcccaagaggagagccggggcctcaagtc	784

RESULT 15
US-08-406-855A-5
Sequence 5, Application US/08406855A
Patent No. 5861309
GENERAL INFORMATION:
APPLICANT: Jonathan A. Bard et al.
TITLE OF INVENTION: DNA Encoding Human Alpha 1 Adrenergic
Sequence 5, Receptors and Uses Thereof
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooper & Dunham LLP
STREET: 1185 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/406,855A
FILING DATE: 21-AUG-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 41337-A-PCT-US/JPW/RDB
TELECOMMUNICATION INFORMATION:

GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.

OM protein - protein search, using sw model

Run on: April 10, 2002, 17:14:57 ; Search time 21.63 Seconds
(without alignments)
433.836 Million cell updates/sec

Title: US-09-380-276A-4
Perfect score: 2255
Sequence: 1 MALKVLEQEKFTFFLLVLL.....LDQSGAIHPATQTSLOEA 417

Scoring table:
BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 212252 seqs, 22503292 residues

Total number of hits satisfying chosen parameters: 212252

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

- Database : Issued Patents AA.*
- 1: /cgn2_6/ptodata/2/iaa/5A_COMB.pap.*
 - 2: /cgn2_6/ptodata/2/iaa/5B_COMB.pap.*
 - 3: /cgn2_6/ptodata/2/iaa/6A_COMB.pap.*
 - 4: /cgn2_6/ptodata/2/iaa/6B_COMB.pap.*
 - 5: /cgn2_6/ptodata/2/iaa/PCTUS_COMB.pap.*
 - 6: /cgn2_6/ptodata/2/iaa/backfiles1.pap.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	869	38.5	210	4	US-09-286-529-3
2	714.5	31.7	151	4	US-09-286-529-4
3	159	7.1	438	1	US-08-097-827-11
4	159	7.1	438	1	US-08-494-574-11
5	150	6.7	206	1	US-08-097-827-7
6	150	6.7	206	1	US-08-494-574-7
7	145	6.4	205	3	US-08-974-022-51
8	145	6.4	205	4	US-08-795-445A-51
9	145	6.4	205	4	US-08-795-447A-51
10	145	6.4	205	4	US-08-974-186-51
11	145	6.4	205	4	US-08-795-446B-51
12	144	6.4	1104	2	US-08-327-832-5
13	144	6.4	1104	2	US-08-828-584-5
14	136	6.0	625	3	US-08-996-139-15
15	136	6.0	625	4	US-08-995-659-15
16	136	6.0	625	4	US-09-215-649A-15
17	134.5	6.0	415	4	US-09-008-353A-6
18	134	5.9	186	1	US-08-089-458B-6
19	134	5.9	307	4	US-08-804-166-4
20	134	5.9	307	4	US-08-910-991-4
21	133.5	5.9	2050	2	US-08-347-594A-2
22	132.5	5.9	197	2	US-08-505-606-1
23	132	5.9	276	4	US-09-041-886-27
24	132	5.9	277	4	US-09-042-785A-10
25	132	5.9	277	4	US-09-006-353A-10
26	131	5.8	139	2	US-08-219-237B-8
27	131	5.8	176	4	US-09-411-722-1

28	130.5	5.8	140	4	US-08-477-347-17	Sequence 17, Appli
29	130.5	5.8	140	4	US-08-476-862-8	Sequence 8, Appli
30	129.5	5.7	336	4	US-08-804-166-8	Sequence 8, Appli
31	129.5	5.7	336	4	US-08-910-991-8	Sequence 8, Appli
32	129	5.7	326	1	US-08-292-549-4	Sequence 4, Appli
33	129	5.7	326	5	PCT-US91-02207-4	Sequence 4, Appli
34	128.5	5.7	1170	1	US-08-313-288B-20	Sequence 20, Appli
35	128	5.7	355	1	US-08-292-549-6	Sequence 6, Appli
36	128	5.7	355	4	US-09-006-353A-14	Sequence 14, Appli
37	127	5.6	419	4	US-08-509-024-7	Sequence 7, Appli
38	127	5.6	419	4	US-09-333-279-7	Sequence 7, Appli
39	126	5.6	148	4	US-09-411-722-2	Sequence 2, Appli
40	124.5	5.5	1111	1	US-08-317-450B-15	Sequence 15, Appli
41	124.5	5.5	1111	4	US-08-800-593-15	Sequence 15, Appli
42	124.5	5.5	1193	1	US-08-317-450B-13	Sequence 13, Appli
43	124.5	5.5	1193	4	US-08-800-593-13	Sequence 13, Appli
44	124.5	5.5	2813	3	US-08-896-449A-2	Sequence 2, Appli
45	124.5	5.5	2813	3	US-09-132-652-2	Sequence 2, Appli

ALIGNMENTS

RESULT 1
US-09-286-529-3
; Sequence 3, Application US/09286529
; Patent No. 6297367
; GENERAL INFORMATION:
; APPLICANT: Catherine Tribouley
; TITLE OF INVENTION: NEW MEMBERS OF TNF AND TNFR FAMILIES
; FILE REFERENCE: 1408.003/200130.439C1
; CURRENT APPLICATION NUMBER: US/09/286,529
; CURRENT FILING DATE: 1999-04-05
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 3
; LENGTH: 210
; TYPE: PRT
; ORGANISM: human
US-09-286-529-3

Query Match 38.5%; Score 869; DB 4; Length 210;
Best Local Similarity 83.7%; Pred. No. 3.9e-73;
Matches 154; Conservative 11; Mismatches 19; Indels 0; Gaps 0;

QY	1	MALKVLEQEKFTFFLLVLLGYSCKVTCTGDCROEFDNRSGNVCPCNQCGMELSK	60
DB	1	MALKVLP L H R T V L F R A I L L H L A C K V S C E T G D C R Q E F D R S G N C V L C K Q C G P G M E L S K	60
QY	61	ECGFGYGEDAQCVTCRLHREFKEDWGFKCKPCLDCAVVRNFQKANC SATSDAICGDCPLG	120
DB	61	ECGFGYGEDAQCVPCPRHFRKEDWGFKCKPCADCALVNRFORANGSHTSDAVCGDCPLG	120
QY	121	FYRKTLVGFQDMCVPCGDP PPPPEPHCASKVNLVKIASTASSPRDTALAAVICSALAT	180
DB	121	FYRKTLVGFQDMCVPCGDP PPPPEPHCTSKVNLVKISSTVSSPRDTAAVAVICSALAT	180

QY	181	VLLA 184
DB	181	VLLA 184

RESULT 2
US-09-286-529-4
; Sequence 4, Application US/09286529
; Patent No. 6297367
; GENERAL INFORMATION:
; APPLICANT: Catherine Tribouley
; TITLE OF INVENTION: NEW MEMBERS OF TNF AND TNFR FAMILIES
; FILE REFERENCE: 1408.003/200130.439C1
; CURRENT APPLICATION NUMBER: US/09/286,529
; CURRENT FILING DATE: 1999-04-05

Db 9 TALLLLG-LTLGVTARRLNCVKHTY--PSGHKC--CRECOPGHGMVNR--DHTRDTLCH 61
QY 74 TCRHLRFKEDWGQKCKPCLDCAVNRFO-KANCATSDAICGDCPLPGFYRKTCLVGFOD 132
Db 62 PCETGYNEAVNYDTCKQCTQCNRHSGSELKQNCPTQDTVC-RCRPGTQPR-----QD 114
QY 133 -----MECPCGDDPPPPYEP-----HCASKVNLVKIATASSPRDTALAIVC---SALA 179
Db 115 SGYKLGVDVCVPC--PPGHFSPGNQACKPWTNCTLSGKQTRHPASDSDAV-CEDRSLLA 171
QY 180 TVLLALLILCVYKQFMEKPSW--SLRSODIOYNGSELSCDLRPOLHEYAHRACCO 236
Db 172 TLL-----WETQRTPTTQSTTVMPTSELP--STPTLVE--PRSC--- 211
QY 237 CRDSVQTCGP 247
Db 212 ---DKTHTCPP 219

RESULT 5

US-08-097-827-7
; Sequence 7, Application US/08097827
; GENERAL INFORMATION:
; APPLICANT: Baum, Peter
; Goodwin, Ray
; Fanslow, William
; Gayle, Richard
; TITLE OF INVENTION: Novel Cytokine Which is a Ligand for
; OX40
; NUMBER OF SEQUENCES: 13
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Immunex Corporation
; STREET: 51 University Street
; CITY: Seattle
; STATE: WA
; COUNTRY: USA
; ZIP: 98101
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/097,827
; FILING DATE: 23-Jul-1993
; CLASSIFICATION: <Unknown>
; ATTORNEY/AGENT INFORMATION:
; NAME: Perkins, Patricia A.
; REGISTRATION NUMBER: 34,693
; REFERENCE/DOCKET NUMBER: 2806
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 206-587-0730
; INFORMATION FOR SEQ ID NO: 7:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 206 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; SEQUENCE DESCRIPTION: SEQ ID NO: 7:
US-08-097-827-7

Query Match 6.7%; Score 150; DB 1; Length 206;
Best Local Similarity 30.1%; Pred. No. 2.4e-06;
Matches 55; Conservative 21; Mismatches 75; Indels 32; Gaps 13;

QY 15 TLLVLGLYSCKVTCGTGCRQOEFRDRSGN-CVPCNOCGPGMELSKGCGFGYGEDAQCV 73
Db 9 TALLLLG-LTLGVTARRLNCVKHTY--PSGHKC--CRECOPGHGMVNR--DHTRDTLCH 61
QY 74 TCRHLRFKEDWGQKCKPCLDCAVNRFO-KANCATSDAICGDCPLPGFYRKTCLVGFOD 132
Db 62 PCETGYNEAVNYDTCKQCTQCNRHSGSELKQNCPTQDTVC-RCRPGTQPR-----QD 114

QY 133 -----MECPCGDDPPPPYEP-----HCASKVNLVKIATASSPRDTALAIVC---SALA 179
Db 115 SGYKLGVDVCVPC--PPGHFSPGNQACKPWTNCTLSGKQTRHPASDSDAV-CEDRSLLA 171
QY 180 TVL 182
Db 172 TLL 174

RESULT 6

US-08-494-574-7
; Sequence 7, Application US/08494574
; Patent No. 5783665
; GENERAL INFORMATION:
; APPLICANT: Baum, Peter
; APPLICANT: Goodwin, Ray
; APPLICANT: Fanslow, William
; APPLICANT: Gayle, Richard
; TITLE OF INVENTION: No. 5783665el Cytokine Which is a Ligand for
; NUMBER OF SEQUENCES: 13
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Immunex Corporation
; STREET: 51 University Street
; CITY: Seattle
; STATE: WA
; COUNTRY: USA
; ZIP: 98101
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/494,574
; FILING DATE: 22-JUN-1995
; CLASSIFICATION: 530
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US/08/097,827
; FILING DATE: 23-JUL-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: Perkins, Patricia A.
; REGISTRATION NUMBER: 34,693
; REFERENCE/DOCKET NUMBER: 2806
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 206-587-0730
; INFORMATION FOR SEQ ID NO: 7:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 206 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-08-494-574-7

Query Match 6.7%; Score 150; DB 1; Length 206;
Best Local Similarity 30.1%; Pred. No. 2.4e-06;
Matches 55; Conservative 21; Mismatches 75; Indels 32; Gaps 13;

QY 15 TLLVLGLYSCKVTCGTGCRQOEFRDRSGN-CVPCNOCGPGMELSKGCGFGYGEDAQCV 73
Db 9 TALLLLG-LTLGVTARRLNCVKHTY--PSGHKC--CRECOPGHGMVNR--DHTRDTLCH 61
QY 74 TCRHLRFKEDWGQKCKPCLDCAVNRFO-KANCATSDAICGDCPLPGFYRKTCLVGFOD 132
Db 62 PCETGYNEAVNYDTCKQCTQCNRHSGSELKQNCPTQDTVC-RCRPGTQPR-----QD 114
QY 133 -----MECPCGDDPPPPYEP-----HCASKVNLVKIATASSPRDTALAIVC---SALA 179
Db 115 SGYKLGVDVCVPC--PPGHFSPGNQACKPWTNCTLSGKQTRHPASDSDAV-CEDRSLLA 171
QY 180 TVL 182

Db 172 TLL 174
!:

RESULT 7

US-08-974-022-51
; Sequence 51, Application US/08974022
; Patent No. 6015938
; GENERAL INFORMATION:
; APPLICANT: Boyle, William J.
; APPLICANT: Lacey, David L.
; APPLICANT: Calzone, Frank J.
; APPLICANT: Chang, Ming-Shi
; TITLE OF INVENTION: OSTEOPROTEGERIN
; NUMBER OF SEQUENCES: 53
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Amgen Inc.
; STREET: 1840 Dehavilland Drive
; CITY: Thousand Oaks
; STATE: California
; COUNTRY: USA
; ZIP: 91320-1789
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/974,022
; FILING DATE: 12-DEC-1995
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/577,788
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Winter, Robert B.
; REFERENCE/DOCKET NUMBER: A-378
; INFORMATION FOR SEQ ID NO: 51:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 205 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-974-022-51

Query Match 6.4%; Score 145; DB 3; Length 205;
Best Local Similarity 28.6%; Pred. No. 6.9e-06;
Matches 54; Conservative 23; Mismatches 76; Indels 36; Gaps 12;

QY 9 QKFTFTLLVLLGLSKVTCTGDCRQOEFRDR--SGN-CVPCNOCGPGMELSKCEGFG 65
Db 6 QQPTAFLLGLSLGVTVKLN-----VKDTYPSGHKC--CRECQPGHGMVSR--D 52

QY 66 YGEDAOCVTCRLHFKEDMGFKCKPCLDCAVNVNRQ--KANCSDATCGDCLPGFYRK 124
Db 53 HTRDTVCHPCPCPGFYNEAVNYDTCKQCTOCNHRSGSELKQNCPTPTDVTVC-QCRPGTQPR 111

QY 125 TKLVGFQDMCEVCPGDPDPPEY-----HCASKVNLV-----KIATASSPRDTALAAVIC- 175
Db 112 QDSSHLKLGVDVCP--PPGHFSPGSNQACKPWTNCTLSGKQIRHPASNSLDT-----VCE 164

QY 176 --SALATVL 182
Db 165 DRSLLATLL 173

RESULT 8
US-08-795-445A-51
; Sequence 51, Application US/08795445A
; Patent No. 6284485
; GENERAL INFORMATION:
; APPLICANT: Boyle, William J.
; APPLICANT: Lacey, David L.
; APPLICANT: Calzone, Frank J.
; APPLICANT: Chang, Ming-Shi
; TITLE OF INVENTION: Osteoprotegerin
; NUMBER OF SEQUENCES: 53
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Amgen Inc.
; STREET: One Amgen Center Drive

; APPLICANT: Boyle, William J.
; APPLICANT: Lacey, David L.
; APPLICANT: Calzone, Frank J.
; APPLICANT: Chang, Ming-Shi
; TITLE OF INVENTION: OSTEOPROTEGERIN
; NUMBER OF SEQUENCES: 53
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Amgen Inc.
; STREET: 1840 Dehavilland Drive
; CITY: Thousand Oaks
; STATE: California
; COUNTRY: USA
; ZIP: 91320-1789
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/795,445A
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/577,788
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Winter, Robert B.
; REFERENCE/DOCKET NUMBER: A-378
; INFORMATION FOR SEQ ID NO: 51:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 205 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-795-445A-51

Query Match 6.4%; Score 145; DB 4; Length 205;
Best Local Similarity 28.6%; Pred. No. 6.9e-06;
Matches 54; Conservative 23; Mismatches 76; Indels 36; Gaps 12;

QY 9 QKFTFTLLVLLGLSKVTCTGDCRQOEFRDR--SGN-CVPCNOCGPGMELSKCEGFG 65
Db 6 QQPTAFLLGLSLGVTVKLN-----VKDTYPSGHKC--CRECQPGHGMVSR--D 52

QY 66 YGEDAOCVTCRLHFKEDMGFKCKPCLDCAVNVNRQ--KANCSDATCGDCLPGFYRK 124
Db 53 HTRDTVCHPCPCPGFYNEAVNYDTCKQCTOCNHRSGSELKQNCPTPTDVTVC-QCRPGTQPR 111

QY 125 TKLVGFQDMCEVCPGDPDPPEY-----HCASKVNLV-----KIATASSPRDTALAAVIC- 175
Db 112 QDSSHLKLGVDVCP--PPGHFSPGSNQACKPWTNCTLSGKQIRHPASNSLDT-----VCE 164

QY 176 --SALATVL 182
Db 165 DRSLLATLL 173

RESULT 9
US-08-795-447A-51
; Sequence 51, Application US/08795447A
; Patent No. 6284728
; GENERAL INFORMATION:
; APPLICANT: Boyle, William J.
; APPLICANT: Lacey, David L.
; APPLICANT: Calzone, Frank J.
; APPLICANT: Chang, Ming-Shi
; TITLE OF INVENTION: Osteoprotegerin
; NUMBER OF SEQUENCES: 53
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Amgen Inc.
; STREET: One Amgen Center Drive

LENGTH: 205 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-795-446B-51

Query Match 6.4%; Score 145; DB 4; Length 205;

Best Local Similarity 28.6%; Pred. No. 6.9e-06;
Matches 54; Conservative 23; Mismatches 76; Indels 36; Gaps 12;

QY 9 QKTEFTLLVLGLYSCKVTCTGCRQOEFRDR--SGN-CVPCNQCGPMELSKGCGFG 65
DB 6 QQTAFLLLGLSLGVTYKLNLC-----VKDTYPSGHKC--CRECQPGHGMVSR--D 52
QY 66 YGEDAQCVTCLRHREKEDMGFKKPCLDCAVNNRFQ-KANGSATSDAICGCLPGFYRK 124
DB 53 HTRDTVCHPCPGFYNEAVNYDTCKQCTQCNHRSGSELKQNTPTEDTVC-QCRPGTQPR 111
QY 125 TKLVGFQDMCEVPCGDPPTPEP---HCASKVNLV---KIATASSPRDTALAAVIC- 175
DB 112 QSSSHKLVDCVPC--PPGHFSGSNQACKPWTNCTLSGKQIRHPASNSLDT-----VCE 164
QY 176 --SALATVL 182
DB 165 DRSLATL 173

RESULT 12

US-08-327-832-5
; Sequence 5, Application US/08327832
; Patent No. 5840832

; GENERAL INFORMATION:

; APPLICANT: Ono, Santa J.

; APPLICANT: Strominger, Jack L.

; TITLE OF INVENTION: Transcription Factor Regulating MHC

; TITLE OF INVENTION: Expression, cDNA and Genomic Clones Encoding Same and
; TITLE OF INVENTION: Retroviral Expression Contracts Thereof

; NUMBER OF SEQUENCES: 16

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: Banner, Birch, McKie & Beckett

; STREET: 1001 G Street, N.W.

; CITY: Washington, D.C.

; STATE: District of Columbia

; COUNTRY: U.S.A.

; ZIP: 20001

; COMPUTER READABLE FORM:

; MEDIUM TYPE: Floppy disk

; COMPUTER: IBM PC compatible

; OPERATING SYSTEM: PC-DOS/MS-DOS

; SOFTWARE: Patent In Release #1.0, Version #1.25

; CURRENT APPLICATION DATA:

; APPLICATION NUMBER: US/08/327,832

; FILING DATE:

; CLASSIFICATION: 530

; ATTORNEY/AGENT INFORMATION:

; NAME: Posorske, Laurence H.

; REGISTRATION NUMBER: 34,698

; REFERENCE/DOCKET NUMBER: 1107,46362

; TELECOMMUNICATION INFORMATION:

; TELEPHONE: 20-2 508-9153

; TELEFAX: 202 508-9299

; INFORMATION FOR SEQ ID NO: 5:

; SEQUENCE CHARACTERISTICS:

; LENGTH: 1104 amino acids

; TYPE: amino acid

; TOPOLOGY: linear

; MOLECULE TYPE: protein

US-08-327-832-5

Query Match

6.4%; Score 144; DB 2; Length 1104;

Best Local Similarity 18.8%; Pred. No. 8.8e-05;
Matches 110; Conservative 59; Mismatches 181; Indels 236; Gaps 27;

QY 24 SKVTCETGDCRQOEFRDRSGNCVPCNQCGPMELSKGCG-----FGYGEDAQC----- 72

DB 440 SCNLLCHPG-----PCPPCPAPMTKTCEGTRHTVRCQGAQSVHCSNPC 484

QY 73 ---VTCRLHREKEDMGFOCKPCLDCAVNNRFQKANGSATS-DAICGDCPLPGFYRKTKLV 128

DB 485 ENILNCGQHQCAELCHGGCQPCQ--IILN--OVYCGSTSRDVLJGTDV-----GKSD 534

QY 129 GFQDMCE-----VPCGD-----PPP-----PYEPHC--ASKYNLVKIASTASS 164

DB 535 GFGDFSLCTGCKDLKGNHSCVCHPQPCQCPRLPOLVRCCPGQTPQLSOLLELSS 594

QY 165 PRDTALAAV-----TC-SALATVLLALLILC-----VYICKRQFMEKK- 201

DB 595 SRKTCMDPVPCGKVGKPLPGGSLDFIHTCEKLCHEGDCGVPVSRVTSVISCRCSEFTKEL 654

QY 202 PMSLRSQDI-----QYNGSELSCLDLDPQ-----LH-----EYA 230

DB 655 PCTSLKSEDAITMCDKRCNKKRLCGRHKCNELCCVDKEHKCPLNCGRLKURGLHRCCEPC 714

QY 231 HRACQO-CRRDSVQT-----CGPVRLPLSMCC-----EE 258

DB 715 HRGNCOTCQWASFDLTHCHGASVIYPPVPCGTRPPECTQTCARVHECDHPVHSHGSEE 774

QY 259 ACS-----PMPATLGGGVHISAASL----- 277

DB 775 KCPCTCTLTQKCMGKHFRSNIPCHLVDISGLPCSATLPCGMHMKQRLCHKGCLVDE 834

QY 278 -----QAR-----NAGPAGEVMPTEFGSLTQSI 300

DB 835 PCKQPCCTTPRADCCHPCMAPHTSSPCPVTAACKAKVLOCECGRRKEMVICSEASTYOR 894

QY 301 CGEFS DANPLMQNMGDNIISCDYPELTGEDIHSLNPELESSTSLDSSNQDLVGGAV 360

DB 895 IAAISMASKITDMLGGS-----VEISKLTIKKEVHQARLECEDECSALERKKR--LAEAF 948

QY 361 PQVSHSENFTAATDLRYNNLTVESASTODALTMRSQLDOESGAI 406

DB 949 HISEDSPFNIRSSGSKFSDSLKEDA--RKDLKFVSDVEKEMETLV 992

RESULT 13

US-08-828-584-5

; Sequence 5, Application US/08828584

; Patent No. 5908762

; GENERAL INFORMATION:

; APPLICANT: Ono, Santa J.

; APPLICANT: Strominger, Jack L.

; TITLE OF INVENTION: Transcription Factor Regulating MHC

; TITLE OF INVENTION: Expression, cDNA and Genomic Clones Encoding Same and
; TITLE OF INVENTION: Retroviral Expression Contracts Thereof

; NUMBER OF SEQUENCES: 16

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: Banner, Birch, McKie & Beckett

; STREET: 1001 G Street, N.W.

; CITY: Washington, D.C.

; STATE: District of Columbia

; COUNTRY: U.S.A.

; ZIP: 20001

; COMPUTER READABLE FORM:

; MEDIUM TYPE: Floppy disk

; COMPUTER: IBM PC compatible

; OPERATING SYSTEM: PC-DOS/MS-DOS

; SOFTWARE: Patent In Release #1.0, Version #1.25

; CURRENT APPLICATION DATA:

; APPLICATION NUMBER: US/08/828,584

; FILING DATE:

; CLASSIFICATION: 435

; ATTORNEY/AGENT INFORMATION:

; NAME: Posorske, Laurence H.

REGISTRATION NUMBER: 34,698
REFERENCE/DOCKET NUMBER: 1107.46362
TELECOMMUNICATION INFORMATION:
TELEPHONE: 20-2 508-9153
TELEFAX: 202 508-9299
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1104 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-828-584-5

Query Match 6.4%; Score 144; DB 2; Length 1104;
Best Local Similarity 18.8%; Pred. No. 8.8e-05;
Matches 110; Conservative 59; Mismatches 181; Indels 236; Gaps 27;

QY 24 SKVVTCTGDCRQEQFRDRSGNVCNCGPGMELSKGCG-----FGYGEDAOC----- 72
DB 440 SCNLLCHPG-----PCPPCPAFMTKTCCGRTTRHTVRCGQAVSVHCSNPC 484
QY 73 ---VTCRLHRFKEDWFOKRCPLDCAVVNRFOKANCATS-DAICGDCPLGCFYRKTILV 128
DB 485 ENILNCGHQHCAELCHGGCCQPCQ--IILN--QVCYCGSTSRDVLGCTDV-----GKSD 534
QY 129 GQDMEC-----VPCGD-----PPHPC--ASKVNLVKIATASS 164
DB 535 GDFGSCSLCTGCKDLKGNHTCSQVCHPOCQCPRLQVRCPCGQTPLSOLLELGS 594
QY 165 PRDTALAAV-----TC-SALATVILALLILC-----VIYKRRQFMKK- 201
DB 595 SRKTCMDPVPCGKVGKPLPCGSLDFIHTCEKLGHEGDCGVPVSRVTSVISCRCSPRTKEL 654
QY 202 PSWSLRSDI-----QYNGSELSCLDLPQ-----LH---EVA 230
DB 655 PCTSLKSEDATPMCDKRCNKKKLGCRHCKNCCVDKHEKPLNCGRLRCLHRCEPC 714
QY 231 HRACQ-CRRDSVQT-----CGVRLLPSCMC-----EE 258
DB 715 HRGNCOTCWOASFDELTHCHGASVIYPPVPCGTRPCTQTCARVHCDHPYHSHSEE 774
QY 259 ACS-----PNPATLGGCVHSAASL----- 277
DB 775 KCPCTFLTKWCMGKHEFRSNIPCHLVDISGLPCSATPLCGMHKCORLCHKHGLVDE 834
QY 278 -----QAR-----NAGPAGEMVPTFFGSILTQSI 300
DB 835 PCKQPCPTPRADCGHPCMAPHTSSPCPVTTACKAKVLOCEGCRKRWICSEASTYQR 894
QY 301 CGEFSADWPLMONGDNISFCDSPYELTGEDIHSLNPELESSTSLDSSQDLVGGAV 360
DB 895 IAAISMASKITDMLGGS-----VEISKLTIKKEVHOARLECEDECSALERKKR--LAEAF 948
QY 361 PVQSHENFTATDLSRYNNTLVESASTQDALTMRSQDOESGAI 406
DB 949 HISEDSPFNIRSSGKFSDSLKEDA--RKOLKFVSDVEKEMETLV 992

RESULT 14
US-08-996-139-15
Sequence 15, Application US/08996139
Patent No. 6017729
GENERAL INFORMATION:
APPLICANT: Anderson, Dirk M.
APPLICANT: Galibert, Laurent
APPLICANT: Maraskovsky, Eugene
TITLE OF INVENTION: Receptor Activator of NF-kappaB
NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
ADDRESSEE: Immunex Corporation, Law Department
STREET: 51 University Street
CITY: Seattle

STATE: WA
COUNTRY: USA
ZIP: 98101
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Power Macintosh
OPERATING SYSTEM: Apple operating System 7.5.5
SOFTWARE: Microsoft Word for Power Macintosh 6.0.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/996,139
FILING DATE: 22 DECEMBER 1997
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 60/064,671
FILING DATE: 14 OCTOBER 1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 08/813,509
FILING DATE: 07 MARCH 1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 08/772,330
FILING DATE: 23 DECEMBER 1996
ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia Anne
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2851-A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
TELEFAX: (206)233-0644
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 625 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-996-139-15

Query Match 6.0%; Score 136; DB 3; Length 625;
Best Local Similarity 22.2%; Pred. No. 0.00022;
Matches 102; Conservative 51; Mismatches 171; Indels 136; Gaps 27;

QY 16 LVLVLGLYSCKVTC-TGDCRQEQFRDRSGNVCNCGPGMELSKGCGFGYGEDAOCVT 74
DB 18 LCVLL--VPLQVTLQVTPCTQERHYEHLGRG--CSRCEPGYLSKSC--TPTSDSVCLP 71
QY 75 CRLHRFKEDWGFQ-KC---KPC-LDCAVV-----NRFQKANCAT-----SDAIC--- 114
DB 72 CGPDEYLDTWNEEDKLLHLKVCDAKALVAVDPGNHTAPRRCACTAGYHWNDSCECCRN 131
QY 115 GDCLPFGYRKTTLVGFQDMECVPC-----GDPPPPY-----BPH----- 148
DB 132 TECAPFGGAQHPQLNKNKDTVCTPCLLGFSDVFSSTDKCPWTNCTLLGKLEAHQGTTS 191
QY 149 ---CASKVNLVKIATASSPRDTALAAVICSALATVLLALLILCVYCKRQFMKKPSWS 205
DB 192 DVCSSMTLRRPPKEAQAAYLPSLI--VLLLFISVVVAAIIFGVYIRKGGKALTANLWN 249
QY 206 -LRSQDIOYNGSELSCLDRLPQLHEYAHRAAC---QCRRDSVOTCGPVRLLPSCMCEACS 261
DB 250 WYNDACSSLSGNKESGDR-----CAGSHSATSSQOEVCCEGILL---MTRKKMV 296
QY 262 PNPATLGGCVHSAAS-----LOARNAGPAGEMVPT-----FFGS 295
DB 297 PEDGAGVCGPVCAAGGPWAEVRDSRTFTLVSEVETQGLSRKIPTDEYTRDPSQSTGS 356
QY 296 LTQSTCGEFSADWPLMONGPM---GGDNISFC-----DSPELT 330
DB 357 LL--LIQCGSKSIPFPQEPLEVGENDSLSQCFGTGTSTVDSGCDFTPEPPTSDMP--V 412
QY 331 GEDIHSLNPELESSTSL-----DSNSSQDLVG--GAVPVQSH 365
DB 413 SPEKH-LIKEIEGDSCLPWWVSSNSTDGYTGSNTFGEH 451

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GenCore version 4.5
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OM nucleic - nucleic search, using sw model

Run on: April 11, 2002, 00:23:10 ; Search time 107.54 Seconds
(without alignments)
2672.499 Million cell updates/sec

Title: US-09-380-276A-5
Perfect score: 1269
Sequence: 1 atggcttaaaagtctact.....ggcagcgactgggttcctcg 1269

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 351203 seqs, 113238999 residues

Total number of hits satisfying chosen parameters: 702406

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Issued_Patents_NA.*
1: /cgn2_6/ptodata/2/ina/5A_COMB.seq.*
2: /cgn2_6/ptodata/2/ina/5B_COMB.seq.*
3: /cgn2_6/ptodata/2/ina/6A_COMB.seq.*
4: /cgn2_6/ptodata/2/ina/6B_COMB.seq.*
5: /cgn2_6/ptodata/2/ina/PCTUS_COMB.seq.*
6: /cgn2_6/ptodata/2/ina/backfiles1.seq.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	435.4	34.3	893	4	US-09-286-529-8
2	314	24.7	623	4	US-09-286-529-9
3	36.4	2.9	1501	1	US-08-722-001-7
4	36.4	2.9	1987	1	US-08-722-001-26
5	36.4	2.9	1997	1	US-08-722-001-27
6	36.4	2.9	2004	1	US-08-722-001-11
7	36.2	2.9	2485	1	US-08-424-424B-1
8	36.2	2.9	2486	5	PCT-US94-05363A-1
9	36	2.8	4136	4	US-09-103-875-2
10	35.6	2.8	1150	4	US-09-372-934-3
11	34.8	2.7	1639	1	US-08-334-698-5
12	34.8	2.7	1639	1	US-08-228-932-5
13	34.8	2.7	1639	1	US-08-468-935-5
14	34.8	2.7	1639	2	US-08-406-855A-5
15	34.8	2.7	1639	2	US-08-722-190-5
16	34.8	2.7	1639	3	US-08-244-354-5
17	34.8	2.7	1639	3	US-09-206-899-5
18	34.8	2.7	1639	5	PCT-US95-04203-5
19	34.6	2.7	5962	6	US36025-5
20	34.6	2.7	7218	1	US-08-232-463-14
21	33.6	2.6	800	2	US-08-416-603-11
22	33.4	2.6	4360	1	US-08-470-350B-1
23	33	2.6	9472	1	US-08-325-547-9
24	32.8	2.6	2230	1	US-08-200-512-1
25	32.6	2.6	1593	2	US-08-524-828-2
26	32.6	2.6	1593	2	US-08-975-114A-2
27	32.6	2.6	1593	3	US-08-849-281A-2

28	32.6	2.6	2247	2	US-08-524-828-1	Sequence 1, Appli
29	32.6	2.6	2247	2	US-08-975-114A-1	Sequence 1, Appli
c 30	32.6	2.6	3891	1	US-08-480-604A-27	Sequence 27, Appl
c 31	32.6	2.6	3891	2	US-08-405-496A-27	Sequence 27, Appl
c 32	32.6	2.6	3891	4	US-08-915-136-27	Sequence 27, Appl
33	32.4	2.6	1167	1	US-07-960-985-1	Sequence 1, Appli
34	32.4	2.6	1167	2	US-08-496-671-1	Sequence 1, Appli
35	32.4	2.6	1280	4	US-09-096-776B-4	Sequence 4, Appli
36	32.4	2.6	1491	4	US-09-082-092-9	Sequence 9, Appli
37	32.4	2.6	1524	4	US-08-840-767-3	Sequence 3, Appli
c 38	32.4	2.6	1690	2	US-08-461-812-3	Sequence 3, Appli
39	32.4	2.6	3083	2	US-08-480-994-36	Sequence 36, Appl
40	32.4	2.6	3083	2	US-08-616-844-36	Sequence 36, Appl
41	32.4	2.6	3083	2	US-08-599-654-36	Sequence 36, Appl
42	32.4	2.6	3083	2	US-08-485-573-36	Sequence 36, Appl
43	32.4	2.6	3083	3	US-08-944-868A-36	Sequence 36, Appl
44	32.4	2.6	3083	3	US-08-944-423A-36	Sequence 36, Appl
45	32.4	2.6	3083	3	US-08-925-743-36	Sequence 36, Appl

ALIGNMENTS

RESULT 1

US-09-286-529-8
; Sequence 8, Application US/09286529
; Patent No. 6297367
; GENERAL INFORMATION:
; APPLICANT: Catharine Tribouley
; TITLE OF INVENTION: NEW MEMBERS OF TNF AND TNFR FAMILIES
; FILE REFERENCE: 1408.003/200130.439C1
; CURRENT APPLICATION NUMBER: US/09/286,529
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 8
; LENGTH: 893
; TYPE: DNA
; ORGANISM: human
US-09-286-529-8

Query Match 34.3%; Score 435.4; DB 4; Length 893;
Best Local Similarity 82.1%; Pred. No. 1.le-125;
Matches 513; Conservative 0; Mismatches 111; Indels 1; Gaps 1;

QY	1	atggcttttaaaagtctactagaacaagaaacacgtttttcactcttttagttacta	60
Db	55	atggcactcaaggctcctcactctacacagagcgtgtctctcgtcgtccattctccta	114
QY	61	ggctatttgtcatgtataaagtgttccctgcacaccagtgtggccaggcaggaattcaag	120
Db	115	ctccacctggcatgtataaagtgtgtgcgaacacggagattgcaggcagcaggaattcaag	174
QY	121	gacggtctggaactgtgttccctgcacaccagtgtggccaggcaggaattgtcttaag	180
Db	175	gacgctatctggaactgtgtcctctgcacacagtgcgaacctggtcgtggttgcacag	234
QY	181	gaatgtggcttcggtctatgggagagatgcacagtgtgtgcgtgccggtcgcacagttc	240
Db	235	gaatgtggcttcggtctatgggagagatgcacagtgtgtgcgtgccggtcgcacagttc	294
QY	241	aaggagactggggtctccagaataatcgaacctgtctgtgactgcgcagtggtgaaaccgc	300
Db	295	aaggagactggggtctccagaataatcgaacctgtgtgactgcgcagtggtgaaaccgc	354
QY	301	tttcagaagcgaattgttcagccaccagtgtatgcacctgcgcgggactgcttgcaggga	360
Db	355	tttcagagggccaactgctcacacaccagtgtatgcctgtcgtcgcgggactgcttgcaggga	414
QY	361	ttttatagaagacgaaactgtcgcgttttcaagacatgagtggtgtccttggagac	420
Db	415	ttttaccggaagaccacaaactggtgtttttcaagacatgagtggtgtccttgcggagac	474

GENERAL INFORMATION:
APPLICANT: Thompson, Wayne J.
TITLE OF INVENTION: ALPHALC ADRENERGIC RECEPTOR ANTAGONISTS
NUMBER OF SEQUENCES: 35
CORRESPONDENCE ADDRESS:
APPLICANT: Nerenberg, Jennie B.
APPLICANT: Lee, Hee-Yoon
APPLICANT: Bell, Ian M.
TITLE OF INVENTION: ALPHALC ADRENERGIC RECEPTOR ANTAGONISTS
NUMBER OF SEQUENCES: 35
CORRESPONDENCE ADDRESS:
ADDRESSEE: Merck & Co., Inc.
STREET: 126 Lincoln Avenue
CITY: Rahway
STATE: New Jersey
COUNTRY: United States of America
ZIP: 07065
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/722,001
FILING DATE:
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/229,276
FILING DATE: 14-APR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Appollina, Mary A.
REGISTRATION NUMBER: 34,087
REFERENCE/DOCKET NUMBER: 19169Y
TELECOMMUNICATION INFORMATION:
TELEPHONE: (908)594-3462
TELEFAX: (908)594-4720
TELEX: 138825
INFORMATION FOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
LENGTH: 1987 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
TOPOLOGY: both
MOLECULE TYPE: cDNA
HYPOTHETICAL: NO
ANTI-SENSE: NO
US-08-722-001-26

Query Match 2.9%; Score 36.4; DB 1; Length 1987;
Best Local Similarity 52.7%; Pred. No. 0.27;
Matches 79; Conservative 0; Mismatches 71; Indels 0; Gaps 0;
QY 474 cgcgtccacgcccctccagccacgggagacggcgctggtcgcttatctgcagcgctct 533
DB 1112 CGAGGACGAGACCATCTGCCATCAACGAGAGCGGGCTCTTCTCGGCTCT 1171
QY 534 ggcacccgtctgctgcccctctctctctctctctctctctctctctctctctctct 593
DB 1172 GGGCTCTTCTACCTGCTTCTGCGCATCATCTCTGCTCATGTACTGCGCGTCTACGTG 1231
QY 594 ggagagaagaaacccagctggtctctctcggtc 623
DB 1232 GGCCAAGAGGGAGAGCGGGGCTCAAGTC 1261

RESULT 5
US-08-722-001-27
Sequence 27, Application US/08722001
Patent No. 5760054
GENERAL INFORMATION:
APPLICANT: Thompson, Wayne J.
APPLICANT: Huff, Joel R.
APPLICANT: Nerenberg, Jennie B.
APPLICANT: Lee, Hee-Yoon

APPLICANT: Bell, Ian M.
TITLE OF INVENTION: ALPHALC ADRENERGIC RECEPTOR ANTAGONISTS
NUMBER OF SEQUENCES: 35
CORRESPONDENCE ADDRESS:
ADDRESSEE: Merck & Co., Inc.
STREET: 126 Lincoln Avenue
CITY: Rahway
STATE: New Jersey
COUNTRY: United States of America
ZIP: 07065
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/722,001
FILING DATE:
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/229,276
FILING DATE: 14-APR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Appollina, Mary A.
REGISTRATION NUMBER: 34,087
REFERENCE/DOCKET NUMBER: 19169Y
TELECOMMUNICATION INFORMATION:
TELEPHONE: (908)594-3462
TELEFAX: (908)594-4720
TELEX: 138825
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
LENGTH: 1997 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
TOPOLOGY: both
MOLECULE TYPE: cDNA
HYPOTHETICAL: NO
ANTI-SENSE: NO
US-08-722-001-27

Query Match 2.9%; Score 36.4; DB 1; Length 1997;
Best Local Similarity 52.7%; Pred. No. 0.27;
Matches 79; Conservative 0; Mismatches 71; Indels 0; Gaps 0;
QY 474 cgcgtccacgcccctccagccacgggagacggcgctggtcgcttatctgcagcgctct 533
DB 1106 CGAGGACGAGACCATCTGCCATCAACGAGAGCGGGCTCTTCTCGGCTCT 1165
QY 534 ggcacccgtctgctgcccctctctctctctctctctctctctctctctctctctct 593
DB 1166 GGGCTCTTCTACCTGCTTCTGCGCATCATCTCTGCTCATGTACTGCGCGTCTACGTG 1225
QY 594 ggagagaagaaacccagctggtctctctcggtc 623
DB 1226 GGCCAAGAGGGAGAGCGGGGCTCAAGTC 1255

RESULT 6
US-08-722-001-11
Sequence 11, Application US/08722001
Patent No. 5760054
GENERAL INFORMATION:
APPLICANT: Thompson, Wayne J.
APPLICANT: Huff, Joel R.
APPLICANT: Nerenberg, Jennie B.
APPLICANT: Lee, Hee-Yoon
APPLICANT: Bell, Ian M.
TITLE OF INVENTION: ALPHALC ADRENERGIC RECEPTOR ANTAGONISTS
NUMBER OF SEQUENCES: 35
CORRESPONDENCE ADDRESS:
ADDRESSEE: Merck & Co., Inc.


```

; ; REFERENCE/DOCKET NUMBER: 325800-1118
; ; TELECOMMUNICATION INFORMATION:
; ; TELEPHONE: 201-994-1700
; ; TELEFAX: 201-994-1744
; ; INFORMATION FOR SEQ ID NO: 1:
; ; SEQUENCE CHARACTERISTICS:
; ; LENGTH: 2486 BASE PAIRS
; ; TYPE: NUCLEIC ACID
; ; STRANDEDNESS: SINGLE
; ; TOPOLOGY: LINEAR
; ; MOLECULE TYPE: gDNA
; ; PCT-US94-05363A-1

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[illegible]

Qy	573	c	573
Db	2144	C	2144

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RESULT      9
US-09-103-875-2/c
; Sequence 2, Application US/09103875A
; Patent No. 6221849
; GENERAL INFORMATION:
; APPLICANT: Szyf, Moshe
; APPLICANT: Bigey, Pascal
; APPLICANT: Ramchandani, Shyam
; TITLE OF INVENTION: DNA METHYLTRANSFERASE GENOMIC SEQUENCES AND ANTISENSE
; TITLE OF INVENTION: OLIGONUCLEOTIDES
; FILE REFERENCE: 106101.194
; CURRENT APPLICATION NUMBER: US/09/103,875A
; CURRENT FILING DATE: 1998-06-24
; EARLIER APPLICATION NUMBER: 60/069,865
; EARLIER FILING DATE: 1997-12-17
; EARLIER APPLICATION NUMBER: 08/866,340
; EARLIER FILING DATE: 1997-05-30
; NUMBER OF SEQ ID NOS: 138
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 2
; LENGTH: 4136
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-103-875-2

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	Query Match	2.8%	Score 36	DB 4	Length 4136	
	Best Local Similarity	49.5%	Pred. No. 0.56			
	Matches 93	Conservative 0	Mismatches 95	Indels 0	Gaps 0	
Qy	689	ccacagagctgctgcagtcgcccgtgactcagtcagacacctgcggccggtgcgt	748			
Db	3917	CCACACGAGCCCTGCTGTCCCTCCTGAGTCGTTCCCCCCCATGGTACCTACCGCC	3858			
Qy	749	tgctccatccatctgctgtgaggagcctgcagcccaacccggcgactcttgattgtg	808			
Db	3857	TGCGGACATCTCGGGCAGCAGATGGCCGGACGGCCAGTCTGGGCACCGGGCTGGGG	3798			
Qy	809	gggtgcattctgcagccagttcttcaggcaagaacgcagggccagccgggagatggtgtgc	868			
Db	3797	CGGTACGCGCCGGATCTTCGGAGGCTTCAGACAGCGCGCGCGCAGCGAGGCGCCCC	3738			

QY 869 cga2tttc 876
I I I I I
Db 3737 GGCTTTC 3730

RESULT 10
US-09-372-934-3
: Sequence 3, Application US/09372934
: Patent No. 6248579
: GENERAL INFORMATION:
: APPLICANT: Stutzman-Engwall, Kim J.
: APPLICANT: McArthur, Hamish
: APPLICANT: Katoh, Yoshihiro
: TITLE OF INVENTION: STREPTOMYCES AVERMITILIS GENE DIRECTING THE RATIO OF
: TITLE OF INVENTION: B2:B1 AVERMECTINS
: FILE REFERENCE: PC10649
: CURRENT APPLICATION NUMBER: US/09/372,934
: CURRENT FILING DATE: 1999-08-12
: EARLIER APPLICATION NUMBER: 60/074,636
: EARLIER FILING DATE: 1998-02-13
: EARLIER APPLICATION NUMBER: PCT/JP99/00130
: EARLIER FILING DATE: 1999-01-25
: NUMBER OF SEQ ID NOS: 25
: SOFTWARE: PatentIn Ver. 2.0
: SEQ ID NO 3
: LENGTH: 1150
: TYPE: DNA
: ORGANISM: Streptomyces hygroscopicus
: FEATURE:
: NAME/KEY: CDS
: LOCATION: (58)..(990)
US-09-372-934-3

	Query Match	2.83;	Score 35.6;	DB 4;	Length 1150;
	Best Local Similarity	57.08;	Pred. No. 0.35;		
	Matches 65;	Conservative 0;	Mismatches 49;	Indels 0;	Gaps 0;
Qy	448	gccagcaaggctcaacctcgtgaagatcgctccacggcctcagccacacggacacggcg	507		
Db	142	gccagacggctaccgcgatcagaagcgctccccggccagggcggtggggactccgag	201		
Qy	508	ctggctgcgcttatctgcagcgctctggccaccgctcctgctggccctgctcacc	561		
Db	202	cgaatccgcgaatgactaatcccatcgtctcccatgataagagacgagtatcctc	255		

RESULT 11
US-08-334-698-5
; Sequence 5, Application US/08334698
; Patent No. 5556753
; GENERAL INFORMATION:
; APPLICANT: Jonathan A. Bard et al.
; TITLE OF INVENTION: DNA Encoding Human Alpha 1 Adrenergic
; TITLE OF INVENTION: Receptors and Uses Thereof
; NUMBER OF SEQUENCES: 6
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: COOPER & DUNHAM
; STREET: 30 Rockefeller Plaza
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10112
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.24
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/334,698
; FILING DATE:
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:

RESULT 13
US-08-468-939-5
; Sequence 5, Application US/08468939
; Patent No. 5714381
; GENERAL INFORMATION:
; APPLICANT: Jonathan A. Bard et al.
; TITLE OF INVENTION: DNA Encoding Human Alpha 1 Adrenoregic
; TITLE OF INVENTION: Receptors and Uses Thereof
; NUMBER OF SEQUENCES: 6
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: COOPER & DUNHAM LLP
; STREET: 1185 Avenue of the Americas
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10036
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.24
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/468,939
; FILING DATE:
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: White, John P.
; REGISTRATION NUMBER: 28,678
; REFERENCE/DOCKET NUMBER: 41337-1B
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212) 278-0400
; TELEFAX: (212) 391-0526
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1639 base pairs
; TYPE: nucleic acid

; STRANDEDNESS: single
; TOPOLOGY: unknown
; MOLECULE TYPE: DNA (genomic)
; HYPOTHETICAL: N
; ANTI-SENSE: N
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 126..1523
; OTHER INFORMATION:
US-08-468-939-5

Query Match 2.7%; Score 34.8; DB 1; Length 1639;
Best Local Similarity 52.0%; Pred. No. 0.76;
Matches 78; Conservative 0; Mismatches 72; Indels 0; Gaps 0;

QY 474 cgcgtccacggcctccagccacgggacacgcgcgtgctgcttctcgcagcgtct 533
DB 635 CGAGGACGAGACCATCTGCCAGATCAACGAGGAGCGGGCTACGTCTTCTCAGCGCT 694
QY 534 ggcacccgtcctgctggtcctctctctgtgtcattctattgtaagagacagtttat 593
DB 695 GGGCTCTTCTACCTGCTCTGGCCATCATCTGCTGCTACTGCTGCTACTGCGGTCTACGTGCT 754
QY 594 ggagagaagaaacccagctgctctctcgcgtc 623
DB 755 GCCCAAGAGGGAGAGCGGGGCTCAAGTC 784

RESULT 14

US-08-406-855A-5
; Sequence 5, Application US/08406855A
; Patent No. 5861309
; GENERAL INFORMATION:
; APPLICANT: Jonathan A. Bard et al.
; TITLE OF INVENTION: DNA Encoding Human Alpha 1 Adrenergic
; TITLE OF INVENTION: Receptors and Uses Thereof
; NUMBER OF SEQUENCES: 23
; CORRESPONDENCE ADDRESS:
; STREET: 1185 Avenue of the Americas
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10036
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/406.855A
; FILING DATE: 21-AUG-1995
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: White, John P.
; REGISTRATION NUMBER: 28,678
; REFERENCE/DOCKET NUMBER: 41337-A-PCT-US/JPW/KDB
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212) 278-0400
; TELEFAX: (212) 391-0526
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1639 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: unknown
; MOLECULE TYPE: DNA (genomic)
; HYPOTHETICAL: N
; ANTI-SENSE: N
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 126..1523

; OTHER INFORMATION:
US-08-406-855A-5

Query Match 2.7%; Score 34.8; DB 2; Length 1639;
Best Local Similarity 52.0%; Pred. No. 0.76;
Matches 78; Conservative 0; Mismatches 72; Indels 0; Gaps 0;

QY 474 cgcgtccacggcctccagccacgggacacgcgcgtgctgcttctcgcagcgtct 533
DB 635 CGAGGACGAGACCATCTGCCAGATCAACGAGGAGCGGGCTACGTCTTCTCAGCGCT 694
QY 534 ggcacccgtcctgctggtcctctctctgtgtcattctattgtaagagacagtttat 593
DB 695 GGGCTCTTCTACCTGCTCTGGCCATCATCTGCTGCTACTGCTGCTACTGCGGTCTACGTGCT 754
QY 594 ggagagaagaaacccagctgctctcgcgtc 623
DB 755 GCCCAAGAGGGAGAGCGGGGCTCAAGTC 784

RESULT 15

US-08-722-190-5
; Sequence 5, Application US/08722190
; Patent No. 5990128
; GENERAL INFORMATION:
; APPLICANT: Charles Gluchowski, Carlos C. Forray, George
; APPLICANT: Chiu, Theresa A. Branche, John M. Wetzel and Paul R. Hartig
; TITLE OF INVENTION: USE OF ALPHA-1C SPECIFIC COMPOUNDS TO
; TITLE OF INVENTION: TREAT BENIGN PROSTATIC HYPERPLASIA
; NUMBER OF SEQUENCES: 6
; CORRESPONDENCE ADDRESS:
; STREET: 1185 Avenue of the Americas
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10036
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.24
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/722.190
; FILING DATE: 4-APR-1995
; CLASSIFICATION: 424
; ATTORNEY/AGENT INFORMATION:
; NAME: White, John P.
; REGISTRATION NUMBER: 28,678
; REFERENCE/DOCKET NUMBER: 41878-D-PCT-JPW/AGL
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212) 278-0400
; TELEFAX: (212) 391-0525
; TELEX:
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1639 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: unknown
; MOLECULE TYPE: DNA (genomic)
; HYPOTHETICAL: N
; ANTI-SENSE: N
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 126..1523
; OTHER INFORMATION:
US-08-722-190-5

Query Match 2.7%; Score 34.8; DB 2; Length 1639;
Best Local Similarity 52.0%; Pred. No. 0.76;

	Matches	78;	Conservative	0;	Mismatches	72;	Indels	0;	Gaps	0;
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Db 635	CGAGGACGGAGACCATCTGCCAGATCAACGAGAGAGCGGGCTACGTGCTTTCTCAGCGCT	694								
QY 534	ggccaccgtcctgtggtgacctctctgtgtcatctattgtaagagagacagtttat	593								
Db 695	GGGCTCCTTCTTACCTGCCCTCTGGCCATCATCCTGGTCATGTACTGCCGGCTCTACGTGGT	754								
QY 594	ggagaagaacccagctggctctctgcggtc	623								
Db 755	GCCCAAGAGGGAGAGCGGGGCTCAAGTC	784								

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GenCore version 4.5
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OM nucleic - nucleic search, using sw model

Run on: April 11, 2002, 00:23:18 ; Search time 107.54 seconds
(without alignments)
3150.559 Million cell updates/sec

Title: US-09-380-276A-7
Perfect score: 1496
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Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 351203 seqs, 113238999 residues

Total number of hits satisfying chosen parameters: 702406

Minimum DB seq length: 0
Maximum DB seq length: 2000000000
Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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2: /cgn2_6/ptodata/2/ina/5B_COMB.seq:*
3: /cgn2_6/ptodata/2/ina/6A_COMB.seq:*
4: /cgn2_6/ptodata/2/ina/6B_COMB.seq:*
5: /cgn2_6/ptodata/2/ina/PTUS_COMB.seq:*
6: /cgn2_6/ptodata/2/ina/backfiles1.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	447.2	29.9	893	4	US-09-286-529-8
2	326.6	21.8	623	4	US-09-286-529-9
3	54.8	3.7	1582	3	US-08-545-196B-10
4	54.8	3.7	1582	3	US-08-545-196B-12
c 5	54.4	3.6	3138	1	US-07-867-106-4
6	53	3.5	2007	3	US-08-747-221B-36
c 7	53	3.5	2007	3	US-08-747-221B-38
8	53	3.5	2007	4	US-09-005-051-36
c 9	53	3.5	2007	4	US-09-005-051-38
10	52.8	3.5	5173	1	US-08-242-677-1
11	52.6	3.5	1420	2	US-08-909-965C-3
12	52.6	3.5	3238	4	US-08-123-934A-5
13	52.6	3.5	3238	5	PCT-US94-10080-5
14	52.4	3.5	3581	2	US-08-738-349-1
15	52	3.5	3709	4	US-09-541-782-7
16	51.8	3.5	3437	3	US-08-860-339-17
17	51.6	3.4	1641	1	US-08-300-903A-8
18	51.2	3.4	991	3	US-08-924-747-25
19	51.2	3.4	991	4	US-09-247-373B-25
20	51.2	3.4	991	4	US-09-296-715-25
21	51.2	3.4	1776	3	US-08-655-352-10
22	50.6	3.4	15124	2	US-08-487-826B-13
23	50.2	3.4	3933	3	US-08-199-776-1
24	50.2	3.4	3933	3	US-08-663-731-1
25	50.2	3.4	3933	3	US-08-879-338-1
26	50.2	3.4	3933	5	PCT-US95-02044-1
27	50	3.3	1046	1	US-08-361-467B-4

28	50	3.3	1046	1	US-08-484-332C-4	Sequence 4, Appli
29	50	3.3	1472	4	US-08-781-420-10	Sequence 10, Appl
c 30	50	3.3	1472	4	US-08-781-420-12	Sequence 12, Appl
31	49.8	3.3	8920	2	US-08-446-855A-1	Sequence 1, Appli
32	49.8	3.3	8920	4	US-09-150-741-1	Sequence 1, Appli
33	49.6	3.3	2700	3	US-09-315-861-1	Sequence 1, Appli
34	49.4	3.3	746	4	US-09-013-810-1	Sequence 1, Appli
35	49.2	3.3	2887	5	PCT-US96-10521-14	Sequence 14, Appli
36	49	3.3	5952	1	US-07-867-106-2	Sequence 2, Appli
37	48.8	3.3	458	1	US-08-524-757-1	Sequence 1, Appli
38	48.8	3.3	2469	4	US-09-111-730-5	Sequence 5, Appli
c 39	48.6	3.2	1582	3	US-08-545-196B-10	Sequence 10, Appl
c 40	48.6	3.2	1582	3	US-08-545-196B-12	Sequence 12, Appl
41	48.6	3.2	2589	4	US-08-569-749-1	Sequence 1, Appli
42	48.6	3.2	2589	5	PCT-US96-112860-1	Sequence 1, Appli
43	48.4	3.2	1393	1	US-08-174-467-18	Sequence 18, Appl
44	48.4	3.2	1393	3	US-08-452-071-18	Sequence 18, Appl
45	48.4	3.2	1490	2	US-08-553-367A-5	Sequence 5, Appli

ALIGNMENTS

RESULT 1
US-09-286-529-8
; Sequence 8, Application US/09286529
; Patent No. 6297367
; GENERAL INFORMATION:
; APPLICANT: Catherine Tribouley
; TITLE OF INVENTION: NEW MEMBERS OF TNF AND TNFR FAMILIES
; FILE REFERENCE: 1408.003/200130.439C1
; CURRENT APPLICATION NUMBER: US/09/286,529
; CURRENT FILING DATE: 1999-04-05
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 8
; LENGTH: 893
; TYPE: DNA
; ORGANISM: human
US-09-286-529-8

Query Match 29.9%; Score 447.2; DB 4; Length 893;
Best Local Similarity 81.6%; Pred. No. 1.5e-101;
Matches 529; Conservative 0; Mismatches 118; Indels 1; Gaps 1;

QY	22	aataatcatattgataaagaagatgctttaaaagtctactagacaagaagaaacgt	81
Db	32	aataaacagtttggtagagccatggcactaaagctcctctacacagacggtgc	91
QY	82	ttttcactcttttagttactaggtctattgtcatgtaaaagtacttgtgaacacaggag	141
Db	92	tcttcgtccattctctcctactccactcgcgtggaagtgagttgcgaacccggag	151
QY	142	actgtagacagaagaattcaggatcggtctgtggaactgtgttcctccgtcaacacagtgtg	201
Db	152	attgcaggcagcaggaaattcaagatcgatctctggaactgtgtcctctgcaaacagtgcg	211
QY	202	gcccaggcatggaattgtctaaagaaatggtctcggctatgggaggaatgcacagtgtg	261
Db	212	gacctggcgtgagttgctccaaggaatggtctcggctatgggaggaatgcacagtgtg	271
QY	262	tgacgtgcgcgctgcacaggttcaaggagactggggctccagaatccaagccctgtc	321
Db	272	tgccctgcaggccgcaccgggttcaaggaaactggggtttccagaagttaagccatgtg	331
QY	322	tggaactgcagtggtgaaccgctttcagaaggaacttgcagccacagtgtatgcga	381
Db	332	cggactgtgcgtggtagaccgctttcagaggcccaactgtcacacaccagtgtatgcgtg	391
QY	382	tctgcgggactgcttcaggatattatagaagacgaacttgcgcttttcaagaca	441
Db	392	tctgcgggactgcttcaggatattatagaagacgaacttgcgcttttcaagaca	451

STREET: PO BOX 747
CITY: FALLS CHURCH
STATE: VA
COUNTRY: USA
ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/545.196B
FILING DATE: 19-OCT-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: FARACI, C. J.
REGISTRATION NUMBER: 32,350
REFERENCE/DOCKET NUMBER: 2121-110P
TELEPHONE: (703) 205-8000
TELEFAX: (703) 205-8050
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 1582 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: CDNA
US-08-545-196B-12

Query Match 3.7%; Score 54.8; DB 3; Length 1582;
Best Local Similarity 67.5%; Pred. NO. 0.00021;
Matches 77; Conservative 0; Mismatches 37; Indels 0; Gaps 0;

Qy 1383 gtttttttttgcaccttaataatttttgcattgtatgttagagtagtgttttaataataat 1442
Db 1417 GCTGTCATGTCAGTGTGTTTCTCTATGTTTAAAGCTATATATAATAAATA 1476
Qy 1443 tccaagtagtttttttaaaactaaaaaataaaaaaataaaaaaataaaaaa 1496
Db 1477 TTTAATTTTTTTTAAAAAATAAAAAAATAAAAAAATAAAAAAATA 1530

RESULT 5
US-07-867-106-4/c
; Sequence 4, Application US/07867106
; Patent No. 5389526
; GENERAL INFORMATION:
; APPLICANT: Slade, Martin B
; APPLICANT: Chang, Andy C M
; APPLICANT: Williams, Keith L
; TITLE OF INVENTION: Improved Plasmid Vectors for Cellular
; TITLE OF INVENTION: Slime Moulds of the Genus Dictyostelium
; NUMBER OF SEQUENCES: 19
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & No. 5389526ris
; STREET: One Liberty Place 46th Floor
; CITY: Philadelphia
; STATE: PA
; COUNTRY: USA
; ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/867.106
FILING DATE: 19920625
PRIOR APPLICATION NUMBER: AU PJ 7187
APPLICATION NUMBER: PCT/AU90/00530

FILING DATE: 02-NOV-1989
ATTORNEY/AGENT INFORMATION:
NAME: Feeney, Joanne Longo
REGISTRATION NUMBER: 35,134
REFERENCE/DOCKET NUMBER: RICE-0002
TELEPHONE: 215-568-3100
TELEFAX: 215-568-3439
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 3138 base pairs
TYPE: NUCLEIC ACID
STRANDEDNESS: single
TOPOLOGY: circular
MOLECULE TYPE: DNA (genomic)
ANTI-SENSE: NO
US-07-867-106-4

Query Match 3.6%; Score 54.4; DB 1; Length 3138;
Best Local Similarity 62.5%; Pred. NO. 0.00035;
Matches 85; Conservative 0; Mismatches 51; Indels 0; Gaps 0;

Qy 1361 aagatttgggggaacctgatgagtttttttttgcattcttaataatttctgtatgttg 1420
Db 2087 AATGTATGTTGGAATTCATTTTATTTTATTTTAAATTTATTTTGTGTTTAAAGAAA 2028
Qy 1421 tagagtgttttaaaataatttcaagtagtttttttaaaactaaaaaataaaaaa 1480
Db 2027 TAAGAAAAAATAAAAAAATAATCTTTTATGCAATCTGAAAAAATAAAAAAATA 1968

Qy 1481 aaaaaaataaaaaa 1496
Db 1967 AAAAAAATAAAAAA 1952

RESULT 6
US-08-747-221B-36
; Sequence 36, Application US/08747221B
; Patent No. 6063610
; GENERAL INFORMATION:
; APPLICANT: Silver, Gary W.
; APPLICANT: Wisniewski, Nancy
; TITLE OF INVENTION: No. 6063610el Carboxylesterase Nucleic Acid
; TITLE OF INVENTION: Molecules, Proteins and Uses Thereof
; NUMBER OF SEQUENCES: 66
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Carol Talkington Verser, Ph.D.
; ADDRESSEE: Heska Corporation
; STREET: 1825 Sharp Point Drive
; CITY: Fort Collins
; STATE: Colorado
; COUNTRY: USA
; ZIP: 80525
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: WordPerfect for Windows, Version 7.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/747.221B
FILING DATE: No. 6063610ember 12, 1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Verser, Carol Talkington
REGISTRATION NUMBER: 37,459
REFERENCE/DOCKET NUMBER: EC-1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 970/493-7272
TELEFAX: 970/484-9505
INFORMATION FOR SEQ ID NO: 36:
SEQUENCE CHARACTERISTICS:
LENGTH: 2007 nucleotides


```
; Patent No. 6291222
; GENERAL INFORMATION:
; APPLICANT: Silver, Gary W.
; APPLICANT: Wisniewski, Nancy
; TITLE OF INVENTION: No. 6291222el Carboxylesterase Nucleic Acid
; TITLE OF INVENTION: Molecules, Proteins and Uses Thereof
; NUMBER OF SEQUENCES: 66
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Carol Talkington Verser, Ph.D.
; ADDRESSEE: Heska Corporation
; STREET: 1825 Sharp Point Drive
; CITY: Fort Collins
; STATE: Colorado
; COUNTRY: USA
; ZIP: 80525
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: Windows 95
; SOFTWARE: WordPerfect for Windows, Version 7.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/005,051
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/747,221
; FILING DATE: No. 6291222ember 12, 1996
; ATTORNEY/AGENT INFORMATION:
; NAME: Verser, Carol Talkington
; REGISTRATION NUMBER: 37,459
; REFERENCE/DOCKET NUMBER: FC-1
; TELEPHONE: 970/493-7272
; TELEFAX: 970/484-9505
; INFORMATION FOR SEQ ID NO: 38:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 2007 nucleotides
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
; US-09-005-051-38

Query Match 3.5%; Score 53; DB 4; Length 2007;
Best Local Similarity 67.9%; Pred. No. 0.00064;
Matches 74; Conservative 0; Mismatches 35; Indels 0; Gaps 0;

Qy 1384 ttttttttgcattttaaataattcttgtagtgtagagtagtattttaaataaatt 1443
      ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 109 TTATTACCATCTTTGTATCATATATTTGCTTTTATTTTTCATTTTTCATTTTTCATTAATA 50

Qy 1444 caagtatattttttaaactaaataaaataaaataaaataaaataaaataaa 1492
      ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 49 TATTGTTTTTATATAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 1

RESULT 10
US-08-242-677-1
; Sequence 1, Application US/08242677
; Patent No. 5677143
; GENERAL INFORMATION:
; APPLICANT: Gaynor, Richard B
; APPLICANT: Wu, Foon W.
; TITLE OF INVENTION: Cellular Nucleic Acid Binding Protein
; TITLE OF INVENTION: and Uses Thereof in regulating Gene Expression and in the
; TITLE OF INVENTION: Treatment of AIDS
; NUMBER OF SEQUENCES: 9
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Arnold, White & Durkee
; STREET: P.O. Box 4433
; CITY: Houston
; STATE: TX
```

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; COUNTRY: USA
; ZIP: 77210
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/242,677
; FILING DATE:
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Mayfield, Denise L.
; REGISTRATION NUMBER: 33,732
; REFERENCE/DOCKET NUMBER: UTSD:401
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 713-787-1400
; TELEFAX: 713-789-2679
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 5173 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 1..4863
; US-08-242-677-1

Query Match 3.5%; Score 52.8; DB 1; Length 5173;
Best Local Similarity 67.0%; Pred. No. 0.0011;
Matches 75; Conservative 0; Mismatches 37; Indels 0; Gaps 0;

Qy 1385 ttttttttgcattttaaataattcttgtagtgtagagtagtattttaaataaattt 1444
      ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 5028 TATTTTCCCTTTAATAAACACTTTTGTAAATGTATCTCTTAAATAAATATTT 5087

Qy 1445 caagtatttttttaaactaaataaaataaaataaaataaaataaaataaa 1496
      ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 5088 TAAGCAATTGTCATATAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 5139

RESULT 11
US-08-909-965C-3
; Sequence 3, Application US/08909965C
; Patent No. 5936078
; GENERAL INFORMATION:
; APPLICANT: Kuga Tetsro
; APPLICANT: Nakagawa Satoshi
; APPLICANT: Sakaki Yoshiyuki
; APPLICANT: Zhao Nanding
; APPLICANT: Hashida Hideji
; TITLE OF INVENTION: NOVEL DNA, NOVEL POLYPEPTIDE
; TITLE OF INVENTION: AND NOVEL ANTIBODY
; NUMBER OF SEQUENCES: 17
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: FITZPATRICK, CELLA, HARPER AND SCINTO
; STREET: 277 Park Avenue
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10172-0194
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/909,965C
; FILING DATE: August 12, 1997
; CLASSIFICATION: 514
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NUMBER OF SEQ ID NOS: 25
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 4
; LENGTH: 151
; TYPE: PRT
; ORGANISM: human
US-09-286-529-4

Query Match 31.3%; Score 714.5; DB 4; Length 151;
Best Local Similarity 82.0%; Pred. No. 4.6e-59;
Matches 123; Conservative 9; Mismatches 17; Indels 1; Gaps 1;

QY 1 MALKVLEQEKFTFTLLVLLGYLSCKVTCTGDC-RQQFRDRSGNCVPCNQCQGMEL 59
DB 1 MALKVLPRLHRTVLFALFLHLLACKVSGTGDSCRQQFRDRSGNCVLCCKQCGPMEL 60

QY 60 KECFGYGEDAQCVTCRLHREFKEDWGFKCPCLDCAVNRFOKANCATSATSDAICGDCLP 119
DB KECFGYGEDAQCVPCRPFRHREFKEDWGFKCPCLDCAVNRFOKANCATSATSDAICGDCLP 120

QY 120 GFYRKTKLVGFQDMCVPCGDP PPPPEPHC 149
DB GFYRKTKLVGFQDMCVPCGDP PPPPEPHC 150

RESULT 3
US-08-097-827-11
; Sequence 11, Application US/08097827
; GENERAL INFORMATION:
; APPLICANT: Baum, Peter
; Goodwin, Ray
; Fanslow, William
; Gayle, Richard
; TITLE OF INVENTION: Novel Cytokine which is a Ligand for
; NUMBER OF SEQUENCES: 13
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Immunex Corporation
; STREET: 51 University Street
; CITY: Seattle
; STATE: WA
; COUNTRY: USA
; ZIP: 98101
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/097,827
; FILING DATE: 23-Jul-1993
; CLASSIFICATION: <Unknown>
; ATTORNEY/AGENT INFORMATION:
; NAME: Perkins, Patricia A.
; REGISTRATION NUMBER: 34,693
; REFERENCE/DOCKET NUMBER: 2806
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 206-587-0730
; INFORMATION FOR SEQ ID NO: 11:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 438 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; SEQUENCE DESCRIPTION: SEQ ID NO: 11:
US-08-097-827-11

Query Match 7.0%; Score 159; DB 1; Length 438;
Best Local Similarity 27.5%; Pred. No. 9.9e-07;
Matches 69; Conservative 31; Mismatches 93; Indels 58; Gaps 18;

QY 15 TLLVLLGYLSCKVTCTGDCRQQFRDRSGN-CVPCNQCQGMELSKCEGFGYGEDAQCV 73
DB TLLVLLGYLSCKVTCTGDCRQQFRDRSGN-CVPCNQCQGMELSKCEGFGYGEDAQCV 74
9 TALLLLG-LTLGVTAARLNCVKHTY--PSGKHC--CRECOPGHGMVNR--DHTRTDLCH 61

QY 74 TCRHREFKEDWGFKCPCLDCAVNRFOKANCATSATSDAICGDCLPGFYRKTKLVGFQD 132
DB TCRHREFKEDWGFKCPCLDCAVNRFOKANCATSATSDAICGDCLPGFYRKTKLVGFQD 133
62 PCETGFYNEAVNYDTCKQCTQCNRHSGSELKNCPTPTQDTVC-RCRPGTQPR-----QD 114

QY 133 -----MECVPCGDP PPPPEPHC-----HCASKVNLVKLTASTASSPRDTALAIVIC---SALA 179
DB MEKVPCGDP PPPPEPHC-----HCASKVNLVKLTASTASSPRDTALAIVIC---SALA 180
115 SGYKLVGDCVPC--PPGHFSPGNQACKPWTNCTLSGKQTRHPASDSDAV-CEDRSLLA 171

QY 180 TVLLALLLVCIVYCKRQFMEKKPSW---SLRSQDIQVNGSELSCLDPRPOLHEVAHRACQ 236
DB TVLLALLLVCIVYCKRQFMEKKPSW---SLRSQDIQVNGSELSCLDPRPOLHEVAHRACQ 237
237 CRDSVOTCGP 247
DB CRDSVOTCGP 248
212 ---DKTHTC PP 219

RESULT 4
US-08-494-574-11
; Sequence 11, Application US/08494574
; Patent No. 5783665
; GENERAL INFORMATION:
; APPLICANT: Baum, Peter
; APPLICANT: Goodwin, Ray
; APPLICANT: Fanslow, William
; APPLICANT: Gayle, Richard
; TITLE OF INVENTION: No. 5783665el Cytokine which is a Ligand for
; NUMBER OF SEQUENCES: 13
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Immunex Corporation
; STREET: 51 University Street
; CITY: Seattle
; STATE: WA
; COUNTRY: USA
; ZIP: 98101
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/494,574
; FILING DATE: 22-JUN-1995
; CLASSIFICATION: 530
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US/08/097,827
; FILING DATE: 23-JUL-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: Perkins, Patricia A.
; REGISTRATION NUMBER: 34,693
; REFERENCE/DOCKET NUMBER: 2806
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 206-587-0730
; INFORMATION FOR SEQ ID NO: 11:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 438 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-08-494-574-11

Query Match 7.0%; Score 159; DB 1; Length 438;
Best Local Similarity 27.5%; Pred. No. 9.9e-07;
Matches 69; Conservative 31; Mismatches 93; Indels 58; Gaps 18;

QY 15 TLLVLLGYLSCKVTCTGDCRQQFRDRSGN-CVPCNQCQGMELSKCEGFGYGEDAQCV 73
DB TLLVLLGYLSCKVTCTGDCRQQFRDRSGN-CVPCNQCQGMELSKCEGFGYGEDAQCV 74

Db 172 TLL 174

RESULT 7

US-08-974-022-51
; Sequence 51, Application US/08974022
; Patent No. 6015938
; GENERAL INFORMATION:
; APPLICANT: Boyle, William J.
; APPLICANT: Lacey, David L.
; APPLICANT: Calzone, Frank J.
; APPLICANT: Chang, Ming-Shi
; TITLE OF INVENTION: OSTEOPROTEGERIN
; NUMBER OF SEQUENCES: 53
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Amgen Inc.
; STREET: 1840 Behavilland Drive
; CITY: Thousand Oaks
; STATE: California
; COUNTRY: USA
; ZIP: 91320-1789
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/974,022
; FILING DATE: 12-DEC-1995
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/577,788
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Winter, Robert B.
; REFERENCE/DOCKET NUMBER: A-378
; INFORMATION FOR SEQ ID NO: 51:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 205 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-974-022-51

Query Match 6.4%; Score 145; DB 3; Length 205;

Best Local Similarity 28.6%; Pred. No. 6.9e-06;

Matches 54; Conservative 23; Mismatches 76; Indels 36; Gaps 12;

QY 9 QEKTFLLVLLGLYLSCKVTCGTGDCRQOEFRDR--SGN-CVPCNOCGPGMELSKCEGFG 65
Db 6 QOPTAFLLGLSLGVTVKLNLC-----VKDTYPSGHKC--CRECPGHGMVSR--D 52
QY 66 YGEDAQCVCRLHRFKEDMGFKCKPCLDCAVVNRFO--KANCSDAICGDCPLCPGYRK 124
Db 53 HTRDTVCHPCPCPGFYNEAVNYDTCKOCTCNHRSGSELKQNCPTPTEDTVC-QCRPGTQPR 111
QY 125 TKLVGFQDMCEVPCGDPPEPPYEP-----HCASKYNLV-----KIASTASSPRDTALAAVIC- 175
Db 112 QDSSHKLGVDCVPC--PPGHFSPGSNQACKPWTNCTLSGKQIRHPASNSLDT-----VCE 164
QY 176 --SALATVL 182
Db 165 DRSLLATLL 173

RESULT 8

US-08-795-445A-51
; Sequence 51, Application US/08795445A
; Patent No. 6284485
; GENERAL INFORMATION:

; APPLICANT: Boyle, William J.
; APPLICANT: Lacey, David L.
; APPLICANT: Calzone, Frank J.
; APPLICANT: Chang, Ming-Shi
; TITLE OF INVENTION: OSTEOPROTEGERIN
; NUMBER OF SEQUENCES: 53
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Amgen Inc.
; STREET: 1840 Behavilland Drive
; CITY: Thousand Oaks
; STATE: California
; COUNTRY: USA
; ZIP: 91320-1789
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/795,445A
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/577,788
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Winter, Robert B.
; REFERENCE/DOCKET NUMBER: A-378
; INFORMATION FOR SEQ ID NO: 51:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 205 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-795-445A-51

Query Match 6.4%; Score 145; DB 4; Length 205;

Best Local Similarity 28.6%; Pred. No. 6.9e-06;

Matches 54; Conservative 23; Mismatches 76; Indels 36; Gaps 12;

QY 9 QEKTFLLVLLGLYLSCKVTCGTGDCRQOEFRDR--SGN-CVPCNOCGPGMELSKCEGFG 65
Db 6 QOPTAFLLGLSLGVTVKLNLC-----VKDTYPSGHKC--CRECPGHGMVSR--D 52
QY 66 YGEDAQCVCRLHRFKEDMGFKCKPCLDCAVVNRFO--KANCSDAICGDCPLCPGYRK 124
Db 53 HTRDTVCHPCPCPGFYNEAVNYDTCKOCTCNHRSGSELKQNCPTPTEDTVC-QCRPGTQPR 111
QY 125 TKLVGFQDMCEVPCGDPPEPPYEP-----HCASKYNLV-----KIASTASSPRDTALAAVIC- 175
Db 112 QDSSHKLGVDCVPC--PPGHFSPGSNQACKPWTNCTLSGKQIRHPASNSLDT-----VCE 164
QY 176 --SALATVL 182
Db 165 DRSLLATLL 173

RESULT 9

US-08-795-447A-51
; Sequence 51, Application US/08795447A
; Patent No. 6284728
; GENERAL INFORMATION:
; APPLICANT: Boyle, William J.
; APPLICANT: Lacey, David L.
; APPLICANT: Calzone, Frank J.
; APPLICANT: Chang, Ming-Shi
; TITLE OF INVENTION: Osteoprotegerin
; NUMBER OF SEQUENCES: 53
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Amgen Inc.
; STREET: One Amgen Center Drive

FILING DATE:

```

RESULT      11
US-08-795-446B-51
; Sequence 51, Application US/08795446B
; Patent No. 6288032
; GENERAL INFORMATION:
; APPLICANT: Boyle, William J.
; APPLICANT: Lacey, David L.
; APPLICANT: Calzone, Frank J.
; APPLICANT: Chang, Ming-Shi
; TITLE OF INVENTION: OSTEOPROTEGERIN
; NUMBER OF SEQUENCES: 53
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Amgen Inc.
; STREET: 1840 Dehavilland Drive
; CITY: Thousand Oaks
; STATE: California
; COUNTRY: USA
; ZIP: 91320-1789
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM pc compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0,
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/795,446
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/577,788
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Winter, Robert B.
; REFERENCE/DOCKET NUMBER: A-378
; INFORMATION FOR SEQ ID NO: 51:
; SEQUENCE CHARACTERISTICS:

```


REGISTRATION NUMBER: 34,698
REFERENCE/DOCKET NUMBER: 1107.46362
TELECOMMUNICATION INFORMATION:
TELEPHONE: 20-2 508-9153
TELEFAX: 202 508-9299
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1104 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-828-584-5

Query Match 6.3%; Score 144; DB 2; Length 1104;
Best Local Similarity 18.8%; Pred. No. 9e-05;
Matches 110; Conservative 59; Mismatches 181; Indels 236; Gaps 27;

QY 24 SKVTCETGDCRQEQFRDRSGNCVPCNCGPGMELSKGCG-----FGYGEDAOC----- 72
DB 440 SCNLLCHPG-----PCPPCPAFMTKTCGGRTRHTVRCGQAVSVHCSNPC 484
QY 73 ---VTCRLHREFKEDGFOCKKPCDCAVNVRFQKANCATS-DAICGDCPLPGFYRKTLV 128
DB 485 ENILNCGQHCAELCHGGCQCPCQ--IILN--QVCYCGSTSRDVLGCTDV-----GKSD 534
QY 129 GFQDMEC-----VPCGD-----PPP-----PYEPHC--ASKVNLVKIATSTASS 164
DB 535 GFGDFSLCTGCKDLKCGNHTSCVCHQPCQCPRLPOLVRCPCGQTPPLSLLELGSS 594
QY 165 PRDTALAAV-----IC-SALATVILALLILC-----VYICKRQFMKK- 201
DB 595 SRKTCMDPVPCGKVGKPLPCGSLDFIHTCEKLCHEGDCGPVSRVTSVISCRCSPRTKEL 654
QY 202 PSWSLRSQDI-----QYNGSELSCLDLDPQ-----LH---EVA 230
DB 655 PCTLSKSEDATFMCDCRKNCKRLGGRHKNEICCVDKHCKPLNCGRLKRLGHLRCEPC 714
QY 231 HRACCO-CRRDSVQT-----CGVRLLPMSCC-----EE 258
DB 715 HRGNCQTCWQASFDLTCGASVIYPPVPCGTRPPECTQTCTCARVHECDHPVYHSHSE 774
QY 259 ACS-----PNPATLGGVHSAASL----- 277
DB 775 KCPPTFTLTKQCMGKHPEFRSNIPCHLVDISGLPCSATLPCGMHKKCORLCHKGECLVDE 834
QY 278 -----QAR-----NAGPAGEMVPTFFGSLTQSI 300
DB 835 PCKQPCITPRADGHPCMAPHTSSPCPVTAACKAKVLOCEGRRKEMVICSEASTYOR 894
QY 301 CGEFSDAWPLMQNPGMDNISCDSYPELTGEDIHSLNPELESSTSLDSNSQDLVGGAV 360
DB 895 IAAISMASKITDMLGGS-----VEISKLTKEVHQARLECDDECSALERKKR--LAEAF 948
QY 361 PVQSHSENFATDLSRYNNTLVESASTODALTMRSLDQESGAI 406
DB 949 HISESDPPEINRSSGKSFSDSLKEDA--RKDLKFVSDVEKMETLV 992

RESULT 14
US-08-996-139-15
Sequence 15, Application US/08996139
Patent No. 6017729
GENERAL INFORMATION:
APPLICANT: Anderson, Dirk M.
APPLICANT: Galibert, Laurent
APPLICANT: Maraskovsky, Eugene
TITLE OF INVENTION: Receptor Activator of NF-kappaB
NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
ADDRESSER: Immunex Corporation, Law Department
STREET: 51 University Street
CITY: Seattle

STATE: WA
COUNTRY: USA
ZIP: 98101
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Power Macintosh
OPERATING SYSTEM: Apple Operating System 7.5.5
SOFTWARE: Microsoft Word for Power Macintosh 6.0.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/996,139
FILING DATE: 22 DECEMBER 1997
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 60/064,671
FILING DATE: 14 OCTOBER 1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 08/813,509
FILING DATE: 07 MARCH 1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 08/772,330
FILING DATE: 23 DECEMBER 1996
ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia Anne
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2851-A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
TELEFAX: (206)233-0644
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 625 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-996-139-15

Query Match 6.0%; Score 136; DB 3; Length 625;
Best Local Similarity 22.2%; Pred. No. 0.00023;
Matches 102; Conservative 51; Mismatches 171; Indels 136; Gaps 27;

QY 16 LVLVLGLYLSCKVTC--TGDRCQEQFRDRSGNCVPCNCGPGMELSKGCGFYGEDAQCVT 74
DB 18 LCVLL--VPLQVTLQVTPPTQERHYEHLGRCC--CSRCEPGKYLSSKC--TPTSDSVCLP 71
QY 75 CRLHREFKEDWGFQ-KC---KPC-LDCAVV-----NRFQKANCST-----SDAIC----- 114
DB 72 CGPDEVLDTWNEEDKCLLHKVCDAGKALVADPGNHTAPRCACCTAGYHWNSDCECCRN 131
QY 115 GDCLPGFYRKTKLVGFQDMECVPC-----GDPPPPY-----EPH----- 148
DB 132 TECAFGGAGHPLQLNKNKTVCCTPCLLGFSDVFSSTDKCPWNTCTLLGKLEAHQGTES 191
QY 149 ---CASKVNLVIATSPASPRDTALAAVICSALATVLLALLILCVYCKRQFMKKPNS 205
DB 192 DVVCSSTWTLRPPRQAQYALPSLI--VULLFISVVVVAIIFGVYRKGKALTANLWN 249
QY 206 -LRSQDIQYNGSELSCLDRLPOLHEYAHRAACC---QCRDVSQVTCGPVRLLPMSMCCSEACS 261
DB 250 WYNDACSSLSGNKESGDR-----CAGSHSATSSQVECEGILL---MTREKMW 296
QY 262 PNPATLGGVHSAAS-----LQARNAGPAGEMVPT-----FFGS 295
DB 297 PEDGAGVCGPVCAAGGPAEVRDSTRFTLVSEVETQGLSRKPTDEYTDPRSPSTGS 356
QY 296 LTQSIGGFSDAWPLMQNPM---GGDNISFC-----DSYPELT 330
DB 357 LL--LIQQGSKSIPPPQEPLEVGENDLSQCFGTGTSTVDSECCDFTTEPPSRDTSMP--V 412
QY 331 GEDIHSLNPELESSTSL-----DSNSSQDLVG-GAVPVQSH 365
DB 413 SPEKH-LTKEIEGDSCLPWWVSSNSTDGYTGSNGTPEGDH 451

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